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(54) Title: COMPOSITIONS AND METHODS FOR POL	LYNUC	CLEOTIDE DELIVERY			
This invention relates to compositions and methods for relates to vectors, targeting ligands, and polycationic agents	s. The p	asing the uptake of polynucleotides into cells. Specifically, the invention polycationic agents are capable of (1) increasing the frequency of uptake if (3) inhibiting serum and/or nuclease degradation of polynucleotides.			

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Compositions and Methods for Polynucleotide Delivery

Description

Field of the Invention

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This invention relates to compositions and methods for increasing the uptake of polynucleotides into cells. Specifically, the invention relates to vectors, targeting ligands, and polycationic agents. The polycationic agents are capable of (1) increasing the frequency of uptake of polynucleotides into a cell, (2) condensing polynucleotides; and (3) inhibiting serum and/or nuclease degradation of polynucleotides.

10 Background of the Invention

Polycations, such as polylysine, have been used to facilitate delivery of nucleic acids to cell interior. Both *in vitro* and *in vivo* applications have taken advantage of this property. See, for example, Gao et al., 1996, Biochem. 35:1027-1036.

Polynucleotides, typically DNA, may be taken into a cell by a receptor-mediated endocytosis pathway, a cellular mechanism which internalizes specific macromolecules. In general, complexes designed to be delivered in this fashion contain nucleic acids encoding the gene of interest and a polycationic agent, which acts as a DNA-binding carrier and both neutralizes the charge on the nucleic acids and condenses it.

Condensation facilitates entry of nucleic acids into cell vesicle systems by simulating a macromolecular structure. For example, polylysine condenses DNA into a toroid or doughnut-like structure. (Wagner et al., 1991, Proc. Natl. Acad. Sci. 88:4255-4259).

Polycations previously utilized for nucleic acid delivery to cell interiors include polylysine, protamines, histones, spermine, spermidine, polyornithine, polyargnine, and putrescine.

All publications mentioned herein are incorporated herein by reference for the purpose of disclosing and describing features of the invention for which the publications are cited in connection with.

Summary of the Invention

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An embodiment of the invention is a vector for expression of polypeptides. The vector of the instant invention comprises: (i) an Epstein Barr Virus (EBV) origin of replication; (ii) a polynucleotide encoding an EBV origin binding protein; (iii) an enhancer; (iv) a promoter; and (v) a terminator. Polynucleotides encoding a desired polypeptide, such as erythropoietin or leptin can be inserted into the vector. Also, ribozyme and antisense polynucleotides can also be inserted into the vector.

One embodiment of the invention is a composition capable of targeting a polynucleotide to a specific cell type. The composition comprises: (i) a lipoprotein; (ii) a polynucleotide binding molecule; and (iii) a polynucleotide.

Another embodiment of the invention is a method of increasing the frequency of uptake of polynucleotides into a cell by contacting a cell with a composition comprising: (i) a lipoprotein, (ii) a polynucleotide binding molecule; and (iii) a polynucleotide.

Yet another embodiment of the invention is a method of increasing the frequency of uptake of polynucleotides into a specific cell type by contacting a population of cells with a composition comprising (i) a lipoprotein, (ii) a polynucleotide binding molecule; and (iii) a polynucleotide.

One embodiment of the invention is a polycationic agent exhibiting a net positive electrical charge at physiological pH with the following formula:

$$Ta - \begin{bmatrix} R_1 & R_2 & O \\ -N - C - C - C - C \end{bmatrix}_n - Tc$$

where Ta and Tc are terminating groups. A preferred subset of these compounds is the set where R_2 is hydrogen. Even more preferred are polymers comprising at least one unnatural amino acid. Also preferred are polymers where R_2 and R_3 are hydrogen and R_1 is not hydrogen, also referred to as poly N-substituted glycines or "poly NSGs."

Another embodiment is a neutral polymer exhibiting no net positive or negative electrical charge at physiological pH with the following formula:

$$Ta - \begin{bmatrix} R_1 & R_2 & O \\ -N - C - C - C \end{bmatrix}_n - Tc$$

where Ta and Tc are terminating groups. A preferred subset of these compounds is the set where R₂ is hydrogen. Even more preferred are polymers comprising at least one unnatural amino acid. Also preferred are polymers where R₂ and R₃ are hydrogen and R₄ is not hydrogen, also referred to as poly N-substituted glycines or "poly NSGs."

The instant polycationic agents and neutral polymers are capable of neutralizing the electrical charge of nucleic acids. A subset of these compounds are capable of (1) condensing the structure of polynucleotides and/ or (2) protecting polynucleotides from serum and/or nuclease degradation.

Yet another embodiment of the invention are polycationic agents and neutral polymers that (1) target binding of nucleic acids to cell surfaces, (2) trigger cell membrane destabilization; (3) exhibit endosome buffering capacity; (4) trigger endocytosis; (5) help trigger the release of polynucleotide/lipid complexes from endosomes or (6) nuclear tropism.

Another embodiment of the invention is a composition comprising a polynucleotide of interest and an effective amount of the polycationic agent to neutralize the charge of the polynucleotide. Optionally, the composition includes a ligand which directs the complex to particular cells expressing a ligand-binding partner, and/or an endosomolytic agent, which serves to cause disruption of the endosome containing the complex.

Another embodiment of the invention is a method of condensing nucleic acids by providing an effective amount of the polycationic agents or neutral polymers of the invention and contacting the agent with the desired polynucleotides.

Also an embodiment of the invention is a method of inhibiting serum and/or nuclease degradation of nucleic acids by providing an effective amount of the the polycationic agents or neutral polymers of the inventions and contacting the agent with the desired nucleic acids.

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Brief Description of the Drawings

Figure 1 is a schematic of a two-step monomer assembly reaction scheme.

Figure 2 is a schematic of a three-step monomer assembly reaction scheme.

Figure 3 is a plasmid map of vector pCMVKmITR-EPI.

Figure 4 is a plasmid map of vector CMVkm2.

Figure 5 is a plasmid map of vector pCMV-KM-cmEPO.

Figure 6 is a plasmid map of vector CMVKmLeptinWt.

Figure 7A illustrates transfection efficiencies for a diverse set of polycationic agents. The polycatonic agents were formulated with DNA at a 2:1, + to - charge ratio and added to either HT1080 (solid bar) or COS (stippled bar) in the presence of 10% serum. Luciferase activity was analyzed 48 hours post-transfection. Total cell protein was measured using a Pierce BCA assay and luciferase activity was normalized against total cell protein.

Figure 7B illustrates the effect of oligomer length on transfection efficiency for polycationic agents having different numbers of the same repeating trimer motif. For both A and B each data point represents the average of 2 experiments.

Figure 8(A-C) shows RZ145-1 peptoid-mediated transfection and transfection mediated by commercially available cationic liposome preparations. RZ145-1 or the indicated lipid was formulated and added to cells in the presence (solid bar) or absence (stippled bar) of 10% serum. Luciferase and total cell protein activity were measured 48 hours after initial transfection. Cells lines are (Figure 8A) 293 human embryonic kidney cells, (Figure 8B) HT1080 human fibrosarcoma cells, and (Figure 8C) NIH03T3 mouse fibroblast cells. Each data point represents the average + strandard error of the mean of three transfections.

Figure 9 illustrates the effect of chloroquine on transfection with RZ145-1 in (A) 293 cells and (B) HT1080 cells. The cells were transfected in the presence (black bar) or absence (stippled bar) of 100 uM chloroquine. Cells were lysed 48 hours post transfection and luciferase activity and total protein content were measured.

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Detailed Description

Definitions

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"Lipoproteins" refers to polypeptides that are associated non-covalently with lipids in the bloodstream and are capable of binding to cellular receptors. Preferably, lipoproteins are those involved with transport and storage of lipids. Such proteins include, for example, chylomicrons, low density lipoprotein (LDL), very low density lipoprotein (VLDL), intermediate density lipoprotein (IDL), and high density lipoprotein (HDL). Also, included in the term are mutants, fragments, or fusions of the naturally occurring lipoproteins. Also, modifications of naturally occurring lipoproteins can be used, such as acetylated LDL.

Mutants, fragments, fusions, or modifications of the naturally occurring lipoproteins are amino acid sequences that exhibit substantial sequence identity to naturally occurring lipoproteins or a fragment thereof. These polypeptides will retain more than about 80% amino acid identity; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polypeptides will exhibit more than about 92% amino acid sequence identity with naturally occurring lipoproteins or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%. All of these polypeptides exhibit receptor binding properties of naturally occurring lipoproteins. Usually, such polypeptides exhibit at least about 20% receptor binding of naturally occurring lipoproteins. More typically, the polypeptides exhibit at least about 40%, even more typically the polypeptides exhibit at least about 60%; even more typically, at least about 70%; even more typically, at least about 80%; even more typically, at least about 85%; even more typically, at least about 90%; even more typically, at least about 95% receptor binding of the naturally occurring lipoproteins.

"Polynucleotide binding molecule" refers to those compounds that associate with polynucleotides, and the association is not sequence specific. For example, such molecules can (1) aid in neutralizing the electrical charge of polynucleotide, or (2) facilitate condensation of nucleotides, or (3) inhibit serum or nuclease degradation.

"Polycationic agent" refers generally to a polymer comprising positively-charged single units, although some non-positively charged units may be present in the polymer. The instant agents exhibit a net positive charge under physiologically relevant pH. Such agents are capable of neutralizing the charge of nucleic acids and can exhibit additional properties, such as condensation and/or serum protection of nucleic acids. Preferably, the agents comprises both amino acids and NSGs as monomeric units; also, preferred are agents comprising NSGs as monomeric units.

"Physiologically relevant pH" varies somewhat between in vitro and in vivo applications. Typically, the physiological pH is at least 5.5; more typically, at least 6.0; even more typically, at least 6.5. Usually, physiologically relevant pH is no more than 8.5; more usually, no more than 8.0; even more usually, no more than 7.5.

"Polynucleotide" or "nucleic acid" refers to DNA, RNA, analogues thereof, peptide-nucleic acids, and DNA or RNA with non-phosphate containing nucleotides. Additionally, these nucleic acids may be single-stranded, double-stranded, or chimeric single- or double-stranded molecules.

The term "oligomer" includes polymers such as poly NSGs, produced by the submonomer process described herein and also in Zuckermann et al., supra. includes polymers, copolymers, and interpolymers of any length. More specifically, oligomers may comprise a single repeating monomer, two alternating monomer units, two or more monomer units randomly and/or deliberately spaced relative to each other. Regardless of the type of polyamide produced, the polyamide of the invention may be produced by the same general procedure which includes repeating a two-step or three step cycle wherein a new monomer unit is added in each cycle until an oligomer of desired length is obtained. The oligomer is preferably 2-100 monomers, more preferably 2-50, or 18-28 monomers or 24 to 48 monomers in length.

The term "frequency of uptake of polynucleotides into a cell" refers to an increase in the amount of polynucleotides actually taken up by a cell relative to the amount actually administered to the cell. The frequency of uptake of polynucleotides into a cell is increased if it is greater than the frequency of uptake of naked polynucleotides. For example, using *in vitro* transfection

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methods, uptake of naked polynucleotides into mammalian cells is not usually detectable over background. Some frequency of uptake, however, can be detected when naked polynucleotides are delivered in vivo. The frequency of uptake in vivo and in vitro depends on the tissue type. The frequency of uptake can be measured by known methods for detecting the presence of polynucleotides, such as Northern, Southern, or Polymerase Chain Reaction (PCR) techniques.

Usually, a composition or compound is capable of increasing the frequency of polynucleotide uptake into a cell if it induces a frequency of uptake that is at least 10% greater than the frequency of naked polynucleotide uptake; more usually, at least 15% greater; even more usually, 20% greater; even more usually, at least 30%; and up to 40% to 100% greater, and even 1,000% and 10,000% greater.

"Naked polynucleotides" refers to polynucleotides that are substantially free from any delivery vehicle that can act to facilitate entry into the cell. For example, polynucleotides are naked when free from any material which promotes transfection, such as liposomal formulations, charged lipids, such as Lipofectin® or precipitating agents such as Ca₁(PO₁)₂.

"Effective amount to increase the frequency of polynucleotide uptake into cells" refers to an amount that induces a frequency of polynucleotide uptake into a cell that is at least 10% greater than the frequency of naked polynucleotide uptake; more usually, at least 15% greater; even more usually, 20% greater; even more usually, at least 30%; even more usually, at least 40%.

"Effective amount to neutralize nucleic acids" refers to the amount used to neutralize at least 10% of the electrical charge of the nucleic acid composition; more preferably; the amount refers to the amount used to neutralize at least 40%; even more preferably, the amount to neutralize 50% of the electrical charge; even more preferably, the amount to neutralize 60% of the electrical charge; even more preferably, the amount to neutralize 70% of the electrical charge; even more preferably, the amount to neutralize 80% of the electrical charge; and most preferably, at least 90% of the electrical charge of the nucleic acid composition of interest.

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"Condensation of nucleic acids" occurs when the polycationic agent that is combined with nucleic acids, neutralizes the electrical charge of the nucleic acids and causes it to assume a reduced structure relative to uncomplexed nucleic acids. Preferably, condensation reduces the structure of nucleic acids to a size that can be internalized by structures present on cell surface membranes. Condensation can be measured by determining the charge of the nucleic acid/polycationic agent by gel electrophoresis, for example. Alternatively, an effective amount to condense nucleic acids can also be measured by the final size of the polycationic agent/nucleic acid complex.

"Effective amount to inhibit serum or nuclease degradation of nucleic acids" refers to the amount used to increase the half-life of the polynucleotide when exposed to serum and/or nucleases by at least 5 minutes as compared the uncomplexed nucleic acids; more preferably, the amount used to inhibit degradation by at least 10 minutes; even more preferably, the amount used to inhibit degradation by at least 30 minutes; even more preferably, the amount used to inhibit degradation by at least 45 minutes; even more preferably, the amount used to inhibit degradation by at least 60 minutes; even more preferably, the amount used to inhibit degradation by at least 90 minutes; and more preferably, the amount used to inhibit degradation by at least 120 minutes.

A composition containing A is "substantially free of" B when at least 85% by weight of the total A+B in the composition is A. Preferably, A comprises at least about 90% by weight of the total of A+B in the composition, more preferably at least about 95% or even 99% by weight.

"Immunogenicity" refers to the ability of a given molecule or a determinant thereof to induce the generation of antibodies with binding capacity to the molecule upon administration in vivo, to induce a cytotoxic response, activate the complement system, induce allergic reactions, and the like. An immune response may be measured by assays that determine the level of specific antibodies in serum, by assays that determine the presence of a serum component that inactivates the polycationic agent/nucleic acid complex or conjugated gene delivery vehicle, or by other assays that measure a specific component or activity of the immune system. As discussed in more detail below, low immunogenicity may be established by these assays. The terms "low immunogenicity," "reduced

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immunogenicity," "lowered immunogenicity" and similar terms are intended to be equivalent terms.

An "origin of replication" is a polynucleotide sequence that initiates and regulates replication of polynucleotides, such as an expression vector. The origin of replication behaves as an autonomous unit of polynucleotide replication within a cell, capable of replication under its own control. With certain origins of replication, an expression vector can be reproduced at a high copy number in the presence of the appropriate proteins within the cell. Examples of origins are the 2μ and autonomously replicating sequences, which are effective in yeast; and the viral T-antigen, effective in COS-7 cells.

General Methods and Detailed Description POLYNUCLEOTIDES

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Polynucleotides used in the instant invention can be used to express desired polypeptides, or can be, in themselves, therapeutic, such as ribozymes or antisense polynucleotides. Such polynucleotides can be used in *in vitro*, *ex vivo*, and *in vivo* applications.

Also, the polynucleotides of the invention can be vectors that express polypeptides, ribozymes, or antisense molecules. Vectors contain at least a promoter to initiate transcription operably linked to the coding region, ribozyme or antisense molecule. Other components that can be included in the vector are, for example: (1) a terminator sequence; (2) a sequence encoding a leader peptide to direct secretion; (3) a selectable marker; and (4) an origin of replication.

An orgin of replication is not necessary. The polynucleotides to be delivered can be either replicating or non-replicating. Other components can be added as desired and convenient.

The polynucleotides and methods of the invention can be utilized with any type of host cell. The choice of promoter, terminator, and other optional elements of an expression vector will depend on the host cell chosen. The invention is not dependent on the host cell selected. Convenience and the desired level of protein expression will dictate the optimal host cell. A variety of hosts for expression are known in the art and available from the American Type Culture Collection (ATCC) (Rockville, Maryland, U.S.A.). Suitable bacterial

hosts suitable include, without limitation: Campylobacter, Bacillus, Escherichia, Lactobacillus, Pseudomonas, Staphylococcus, and Streptococcus. Yeast hosts from the following genera may be utilized: Candida, Hansenula, Kluyveromyces, Pichia, Saccharomyces, Schizosaccharomyces, and Yarrowia. Aedes aegypti, Bombyx mori, Drosophila melanogaster, and Spodoptera frugiperda (PCT Patent Publication No. WO 89/046699; Carbonell et al., 1985, J. Virol. 56:153; Wright, 1986, Nature 321:718; Smith et al., 1983, Mol. Cell. Biol. 3:2156; and see generally, Fraser et al., 1989, In Vitro Cell. Dev. Biol. 25:225).

Useful mammalian cell types for *in vitro* applications include for example, those cell lines available from the American Type Culture Collection (ATCC), Chinese hamster ovary (CHO) cells, HeLa cells, baby hamster kidney (BHK) cells, monkey kidney cells (COS), human hepatocellular carcinoma cells (e.g., Hep G2), human embryonic kidney cells, baby hamster kidney cells, mouse sertoli cells, canine kidney cells, buffalo rat liver cells, human lung cells, human liver cells, mouse mammary tumor cells, as well as others.

Useful cell types for *in vivo* or *ex vivo* applications include, without limitation, any tissue type, such as muscle, skin, brain, lung, liter, spleen, blood, bone marrow, thymus, heart, lymph, bone, cartilage, pancreas, kidney, gall bladder, stomach, intestine, testis, ovary, uterus, rectum, nervous system, eye, gland, and connective tissue.

A. In vitro and Ex vivo Vectors

The polynucleotides encoding the desired polypeptides or ribozymes, or antisense polynucleotides can be transcribed and/or translated using the following promoters and enhancers as examples. The examples include, without limitation: the 422(aP2) gene and the stearoyl-CoA desaturase 1 (SCD1) gene, which contains suitable adipocyte-specific promoters, as described in Christy et al., 1989, Genes Dev. 3:1323-1335. Synthetic non-natural promoters or hybrid promoters can also be used herein. For example, a T7T7/T7 promoter can be constructed and used, in accordance with Chen et al., 1994, Nucleic Acids Res. 22:2114-2120, where the T7 polymerase is under the regulatory control of its own promoter and drives the transcription of a polynucleotide sequence, which is

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placed under the control of another T7 promoter. The primary determinant for the fat-specific expression is an enhancer located at about >5 kb upstream of the transcriptional start site, as described in Ross et al., 1990, Proc. Natl. Acad. Sci. USA.87:9590-9594 and Graves et al., 1991, Genes Dev. 5:428-437. Also suitable for use herein is the gene for the CCAAT/enhancer-binding protein C/EBPα, which is highly expressed when 3T3-L1 adioblast commit to the differentiation pathway and in mature post-mitotic adipocytes, as described in Birkenmeier et al., 1989, Gene Dev. 3:1146-1156. The recently isolated transcription factor PPARγ2, expressed exclusively in adipocyte tissues, as described in Tontonoz et al., 1994, Cell 79:1147-1156, can also be used herein.

Typical promoters for mammalian cell expression include the SV40 early promoter, the CMV promoter, the mouse mammary tumor virus LTR promoter, the adenovirus major late promoter (Ad MLP), and the herpes simplex virus promoter, among others. Other non-viral promoters, such as a promoter derived from the murine metallothionein gene, will also find use in mammalian constructs. Expression may be either constitutive or regulated (inducible), depending on the promoter. Typically, transcription termination and polyadenylation sequences will also be present, located 3' to the translation stop codon. Preferably, a sequence for optimization of initiation of translation, located 5' to the coding sequence, is also present. Examples of transcription terminator/polyadenylation signals include those derived from SV40, as described in Sambrook et al., 1989, "Molecular Cloning, A Laboratory Manual," second edition, Cold Spring Harbor Press, Cold Spring Harbor, New York. Introns, containing splice donor and acceptor sites, may also be designed into the constructs of the present invention.

Enhancer elements can also be used herein to increase expression levels of the mammalian constructs. Examples include the SV40 early gene enhancer, as described in Dijkema et al., 1985, EMBO J. 4:761, and the enhancer/promoter derived from the long terminal repeat (LTR) of the Rous Sarcoma Virus, as described in Gorman et al., 1982b, Proc. Natl. Acad. Sci. USA 79:6777, and human cytomegalovirus, as described in Boshart et al., 1985, Cell 41:521. A leader sequence can also be present which includes a sequence encoding a signal peptide, to provide for the secretion of the foreign protein in mammalian cells. Preferably, there are processing sites encoded between the leader fragment and the gene of interest such that the leader sequence can be cleaved either in vivo or in vitro.

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Other regulatory regions from viruses can be included in the polynucleotides of the instant invention to increase transcription and translation levels or increase the duration of transcription and translation. For example, the long terminal repeats of HIV can be included. Alternatively, the inverted terminal repeats of the Epstein Barr Virus can be used.

There exist expression vectors that provide for the transient expression in mammalian cells of DNA encoding the target polypeptide. In general, transient expression involves the use of an expression vector that is able to replicate efficiently in a host cell, such that the host cell accumulates many copies of the expression vector and, in turn, synthesizes high levels of a desired polypeptide encoded by the expression vector. Transient expression systems, comprising a suitable expression vector and a host cell, allow for the convenient positive identification of polypeptides encoded by cloned DNAs, as well as for the rapid screening of such polypeptides for desired biological or physiological properties. Thus, transient expression systems are particularly useful for purposes of identifying analogs and variants of the target polypeptide that have target polypeptide-like activity.

B. In vivo Vectors

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For delivery using viral vectors, any of a number of viral vectors can be used, as described in Jolly, 1994, Cancer Gene Therapy 1:1-64. For example, the coding sequence of a desired polypeptide or ribozymes or antisense molecules can be inserted into plasmids designed for transcription and/or translation in retroviral vectors, as described in Kimura et al., 1994, Human Gene Therapy 5:845-852, adenoviral vectors, as described in Connelly et al., 1995, Human Gene Therapy 6:185-193, adeno-associated viral vectors, as described in Kaplitt et al., 1994, Nature Genetics 6:148-153 and sindbis vectors. Promoters that are suitable for use with these vectors include the Moloney retroviral LTR, CMV promoter and the mouse albumin promoter. Replication competent free virus can be produced and injected directly into the animal or humans or by transduction of an autologous cell ex vivo, followed by injection in vivo as described in Zatloukal et al., 1994, Proc. Natl. Acad. Sci. USA 91:5148-5152.

The polynucleotide encoding a desired polypeptide or ribozyme or antisense polynucleotide can also be inserted into plasmid for expression of the desired polypeptide *in vivo*. For *in vivo* therapy, the coding sequence can be delivered by direct injection into tissue, or via oral administration as an aerosol. Promoters suitable for use in this manner include endogenous and heterologous promoters such as CMV. Further, a synthetic T7T7/T7 promoter can be constructed in accordance with Chen et al., 1994, Nucleic Acids Res. 22:2114-2120, where the T7 polymerase is under the regulatory control of its own promoter and drives the transcription of polynucleotide sequence, which is also placed under the control of a T7 promoter. The polynucleotide can be injected in a formulation that can stablize the coding sequence and facilitate transduction thereof into cells and/or provide targeting, as described in Zhu et al., 1993, Science 261:209-211.

Expression of the coding sequence of a desired polypeptide or replication of a ribozyme or antisense polynucleotide *in vivo* upon delivery for gene therapy purposes by either viral or non-viral vectors can be regulated for maximal efficacy and safety by use of regulated gene expression promoters as described in Gossen et al., 1992, Proc. Natl. Acad. Sci. USA 89:5547-5551. For example, the polynucleotide transcription and/or translation can be regulated by tetracycline responsive promoters. These promoters can be regulated in a positive or negative fashion by treatment with the regulator molecule.

For non-viral delivery of the coding sequence of the desired polypeptide, the sequence can be inserted into conventional vectors that contain conventional control sequences for high level expression.

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C. Preferred Vector

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A preferred vector comprises: (1) an (EBV) origin of replication or a BKV (BK virus), a parvovirus, origin of replication; (2) a coding region for an EBV or BKV origin binding protein; (3) at least one inverted terminal repeat; (4) a promoter; (5) an enhancer; (6) a terminator; (7) optionally, a selectable marker.

Preferably, the orgin of replication is EBV ori p; more preferably, nucleotides 2623 to 4559 of SEQ ID NO:1 are utilized. The sequence is obtainable from vector pCEP4, commercially available from Invitrogen, San Diego, California, USA.

Preferably, the coding region encodes the EBV nuclear antigen A, which binds to EBV ori p; more preferably, the polynucleotide sequence is nucleotides 14 to 2594 of SEQ ID NO:1 are utilized. The sequence is obtainable from vector pCEP4, commercially available from Invitrogen, San Diego, California, USA.

Fragments and mutants of the preferred origin and binding protein capable of initiating replication of the vector in the desired host cell can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 14 to 2594 or 2623 to 4559 of SEQ ID NO:1 or fragment thereof; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 14 to 2594 or 2623 to 4559 of SEQ ID NO:1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%.

Preferably, the inverted terminal repeats are those sequences found in adenovirus (AV) or adeno-associated virus (AAV); more preferably, the inverted terminal repeats are those found in AAV; even more preferably, the polynucleotide sequence is 4938 to 5104 or 7189 to 7355 of SEQ ID NO: 1. The sequence of AAV is described in Samulski et al., 1987, J. Virol. 61:3096-3101.

Fragments and mutants of the preferred inverted terminal repeat capable of initiating replication of the vector in the desired host cell can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 4938 to 5104 or 7189 to 7355 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at

least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 4938 to 5104 or 7189 to 7355 of SEQ ID NO: lor fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%.

Preferably, the cytomegalovirus enhancer/promoter is utilized; more preferably, the CMV promoter sequence is nucleotide sequence 5112 to 6734 of SEQ ID NO:1.

Mutants and fragments of the preferred enhancer and promoter capable of initiating transcription and/or translation can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 5112 to 6734 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 5112 to 6734 of SEQ ID NO: 1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 99%.

A preferred terminator is the bovine growth hormone poly A sequence; more preferably, the polynucleotide sequence is nucleotide 6818 to 7050 of SEQ ID NO:1.

Mutants and fragments of the preferred terminator capable of terminating transcription and/or translation can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 6818 to 7050 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 6818 to 7050 of SEQ ID NO: 1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%.

The sequence of the preferred vector is shown in SEQ ID NO:1. Polynucleotides encoding polypeptides, such as erythropoeitin or leptin, and ribozymes and antisense polynucleotides can be inserted into the vector.

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D. Examples of Coding Regions, Ribozymes, and Antisense Molecules

The following are examples of coding regions, ribozymes, and antisense molecules that can be used to treat various indications in mammals. The nucleotide sequence of the genes of interest can be found, for example, in publically available databases, such as Genbank. Polynucleotides to be delivered can be used to treat viral infections or chronic pathogen infection.

1. Hemophilia

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Gene replacement by *in vivo* delivery of polynucleotides can be effective in treating hemophilia. The following are examples of polypeptides that can be encoded by the polynucleotides to be delivered: Factor VIII:C, mutants of Factor VIII:C, preferably those that are uncleavable. Also, useful to treat hemophilia are ribozyme and antisense polynucleotides as inhibitors of Tissue Factor Plasminogen Inhibitor (TFPI).

The routes of delivery for treating hemophilia include, for example, intravenous/intrahepatic injection, ex vivo transduction of stem cells or lymphocytes using retroviral vectors.

2. Treatment of Graft Versus Host Disease

In vivo delivery of polynucleotides encoding prodrugs can be used for direct ablation to treat graft versus host disease in, for example, leukemia bone marrow transplantation. Herpes thymidine kinase in conjunction with gancyclovir can be utilized for this purpose. Other examples of prodrugs are described in the cancer section.

The routes of delivery for treating graft versus host disease include, for example, ex vivo transduction of T-lymphocytes using retroviral vectors.

3. Vaccines

In vivo delivery of polynucleotides encoding a desired antigen can be utilized to induce an immune response. This response can include both cellular and humoral response. This type of vaccine can be used to treat cancer as well as infectious diseases. Further, such treatment can be either prophylactic or therpeutic immunotherapy.

Examples of infectious diseases include, Human Immunodeficiency Virus (HIV), Hepatitis A, B, C, etc., (HAV, HBV, HCV, etc.), Human Papiloma Virus (HPV), cytomegalovirus (CMV), herpes simplex 1 and 2 (HSV), etc. Preferred antigens include non-structural proteins 3, 4a, and 5b (NS3, NS4, and NS5b) of HCV; gB2 and gD2 of HSV; env and rev proteins of HIV.

Also, cancer antigens can be used in vaccines, for both therapeutic and prophylactic purposes.

The antigens can be presented in the context of Class I major histocompatibility antigens, or to induce a cellular cytotoxic T cell response, or to induce a humoral response comprising the synthesis of antibodies.

In addition, an antisense or ribozyme target to a immune suppressive molecule, IL-10, TGF-β, and CTLA-4, for example, can be useful to be administered with a vaccine.

The routes of delivery for vaccines include, for example, intramuscular injection, dendritic cell-based immunization, or oral immunization by both viral and non-viral vectors.

4. Diabetes Mellitus

Diabetes is another indication that can be treated by *in vivo* delivery of a replacement gene. The following are examples of useful polypeptides to be encoded by the replacement gene: insulin, insulin-like growth factor I and II (IGF-I and II).

Also useful for treating diabetes are polynucleotides encoding IAS-L, found on the surface of B cells in the pancreas, to protect the cells from immune destruction.

The routes of delivery for treating diabetes include, for example, liverdirected, parotid- directed, pancreas-directed, salivary gland-directed using both viral and non-viral vectors.

5. Hyperlipidemia

Hyperlipidemia can be treated by *in vivo* delivery of the following

polynucleotides encoding apoproteins or lipoprotein receptors. A more extensive description of lipoproteins and apoproteins is provided below.

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The routes of delivery for treating hyperlipidemia include, for example, liver-directed intravenous administration by both viral and non-viral vectors.

6. Myocardial Ischemia or Infarction

The following are examples of polynucleotides that are useful, when delivered *in vivo*, to treat myocardial ischemia or infarction:

polynucleotides encoding basic fibroblast growth factor (bFGF), fibroblast growth factor 5 (FGF-5) and IGF-I.

The routes of delivery for treating myocardial ischemia or infarction include, for example, intrapericardial delivery of viral vector or non-viral vectors.

7. Bowel Disease

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The following are examples of polynucleotides that can be delivered in vivo to treat bowel disease:

- (i) ribozymes or antisense polynucleotides as inhibitors of macrophage/inflammatory cell recruitment or activation, such as NFκB;
 - (ii) ribozymes or antisense polynucleotides to act as anti-apoptotic agents, such as inhibitors of interleukin 1b converting enzyme family;
 - (iii) polynucleotides encoding complement blockers, such as decay accelarting factor (DAF), membrane cofactor protein (MCP); and the fusions of DAF and MCP also known as CAB-2;
 - (iv) cyclooxygenase inhibitors;
 - (v) anti-proliferative agents, such as, ribozymes, antisense oligonucleotides, antibodies, protein, or peptides against c-myb, ras/raf, PI3 kinase, cyclins;
- 25 (vi) polynucleotides encoding suicide proteins/genes, such as, herpes thymidine kinase;
 - (vii) polynucleotides encoding replacement genes or proteins which maybe deficient or down regulated during the devleopment of inflammatory bowel disease.
- 30 (viii) polynucleotides encoding IkB.

8. Prostate Cancer and Benign Prostatic Hyperplasia

The following polynucleotides can be delivered to treat prostate cancer and benign prostatic hyperplasia:

- (i) a polynucleotide encoding a pro-apoptotic agent, including for example, fas, fas ligand, fadd, fap-1, tradd, faf, rip, reaper, apoptin, interleukin-2 converting enzyme;
- (ii) a polynucleotide encoding an anti-angiogenic agent, including, for example, bFGF soluble receptor and fragments, angiostatin, transforming growth factor- β (TGF- β), interferon- α (IFN α), proliferin-related protein, a urokinase plasminogen activator receptor antagonist, platelet factor 4 (PF4), thrombospondin, a tissue inhibitor of metalloproteinase, and prolactin;
- (iii) a polynucleotide encoding a immunomodulating agent including, for example, interleukin-2 (IL-2), IFNα, IFNβ, IFNγ, granulocyte macrophage-colony stimulating factor (GM-CSF), and macrophage-colony stimulating factor (M-CSF);
- (iv) a ribozyme or antisense polynucleotide as an antiproliferative agent including, for example, an inhibitor of a signal transduction pathway, for example, an inhibitor of a signal transduction pathway mediated by myb, ras, ras superfamily, raf, phosphoinositol (PI3-kinase), a phosphotyrosine binding (PTB) domain, a SRC homology-2 (SH2) domain, a SRC homology-3 (SH3) domain, a plextrin homology (PH) domain, JUN kinase, and a stress activated kinase, signaling inositol phosphatases; and an inhibitor of a cyclin;
- (v) a ribozyme or antisense polynucleotide as an inhibitor of a growth factor or inhibitor of a receptor of a growth factor, including, for example, epidermal growth factor (EGF), TGF-α, FGF, TGF-β, platelet derived growth factor (PDGF), keratinocyte growth factor (KGF), or any prostate cell specific growth factor;
- (vi) a polynucleotide encoding a tumor suppressor gene or a gene downregulated during the onset of a hyperplastic condition in the prostate; and
- (vii) an antisense or ribozyme target to a immune suppressive molecule, IL-10, TGF-β, and CTLA-4, for example.

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9. Anemia, Leukopenia, and Thrombocytopenia

Anemia can be treated by *in vivo* delivery of a polynucleotide encoding erythropoietin, GM-CSF-, G-CSF, M-CSF, and thrmobopoietin, for example. Examples of delivery routes for this indication include without limitation: livertargeted intravenous administration of viral vectors and non-viral vectors. See the Examples below.

10. Cardiomyopathy

The following are examples of polynucleotides that can be delivered in vivo to treat cardiomyopathy: polynucleotides encoding, IGF-1, L-amino acid decarboxylas, inhibitors of β adrenergic receptor kinases (BARK), troponin, and β adrenergic receptors.

Examples of delivery routes for this indication include, without limitation, pericardial expression of IGF-1, and for the other genes, intramycardial injection or myocardial trageting via intracoronary injection or intrapericardial administration of viral vectors or non-viral vectors.

11. Rheumatoid Arthritis

The following are examples of polynucleotides that can be delivered in vivo to treat rheumatoid arthritis, polynucleotides encoding a prodrug, such as herpes thymidine kinase, MMP inhibitors, fas, and pro-apoptotic proteins, described above, and interleukin-1 receptor A, interleukin-10, IkB.

Also, antisense and ribozyme polynucleotides as inhibitors of NFkB. Examples of delivery routes for this indication include, without limitation, intraarticular injection of viral and non-viral vectors.

12. Osteoarthritis and Psoriasis

The following are examples of polynucleotides that can be delivered in vivo to treat osteoarthritis and psoriasis: polynucleotides encoding IGF-1; ribozyme and antisense polynucleotides as inhibitors of metalloproteinase inhibitors.

Also, the following are examples of polynucleotides that can be delivered in vivo to treat osteoarthritis and psoriasis, polynucleotides encoding a prodrug.

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such as herpes thymidine kinase, MMP inhibitors, fas, and pro-apoptotic proteins, described above, and interleukin-1 receptor A, interleukin-10, IkB.

Also, antisense and ribozyme polynucleotides as inhibitors of NFkB.

Examples of delivery routes for this indication include, without limitation, intraarticular injection.

13. Restenosis

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The following are examples of polynucleotides that can be delivered in vivo to treat restenosis:

- (i) polynucleotides encoding a prodrug, such as thymidine kinase, other examples are described in the cancer section;
- (ii) polynucleotides encoding tissue factor plasminogen inhibitor (TFPI);
 - (iii) polynucleotides encoding c-myb rbz, c-ras rbz,
 - (iv) polynucleotides encoding pro-apoptotic agents, described above;
 - (v) polynucleotides encoding IkB.

Examples of delivery routes for this indication include, without limitation, intracoronary delivery of viral and non-viral vectors.

14. Cancer

The gene delivery vectors of the invention are useful in delivering therapeutic genes for treatment of hyperproliferative disorders, including malignancy, for treatment of infectious disease and for treatment of inflammatory diseases, including autoimmune disease. For instance, the gene therapy vectors can be used to express cytokines or proteins that convert an inactive or partially active prodrug into an active drug. In many cases, conversion of the prodrug into its active form results in a compound with cytolytic activity.

a. Prodrug converting enzymes

A number of "suicide genes" which encode different proteins useful in prodrug conversion can be used in the instant invention. For instance, nucleoside kinases such as thymidine kinase are particularly useful. In particular, the HSV-

TK system has important advantages for anti-tumor cell therapy. See PCT publication number WO 91/02805 entitled "Recombinant Retroviruses Delivering Vector Constructs to Target Cells" and PCT publication number WO 95/14014091 entitled "Compositions and Methods for Utilizing Conditionally Lethal Genes" for a description of treatment of cancer and other diseases by gene 5 delivery vectors expressing thymidine kinase and other prodrug converting enzymes. HSV-TK transduced tumor cells can mediate a significant bystander killing effect on untransduced neighboring cells in vitro and in vivo (Moolten et al., supra., Freeman et al., 1993, Cancer Res. 53:5274), most commonly as a 10 result of transfer to the toxic ganciclovir metabolite, GCV triphosphate, between adjacent cells through intercellular gap junctions (Bi et al., 1993, Human Gene Therap. 4:725). Endothelial cells in capillary walls are connected by gap junctions, so a dramatic bystander effect created by GCV-triphosphate transfer between neighboring endothelial cells and the massive amplification effects of 15 the clotting cascade and the tumor to endothelial cell ratio could ensue (Denekamp et al., 1986, <u>Cancer Topics</u> 6:6; Denekamp et al., 1984, Prog. Appl. Microcir. 4:28). Recent evidence suggests that the occasional transduction of tumor endothelial cells during intralesional therapy with HSV-TK retroviral vectors may account for a significant component of the antitumor activity of the vectors (Ram et al., 1994, J. Neurosurg. 81:256). In addition, the suicide gene is 20 only conditionally cytotoxic to the target cells (i.e. only when GCV is given). Consequently, an ex vivo administration method can be be utilized. For example, in this type of protocol, endothelial cells may be isolated from tumor biopsies (Medzelewski et al., 1994, Cancer Res. 54:336), induced to proliferate with 25 appropriate mitogens (Ferrara et al., supra.) and transduced with TK in vitro. Transplanted EC become incorporated into the neovasculature in days to weeks after intratumoral injection (Lal et al., 1994, Cancer Gene Therap.1:322), so GCV treatment would follow a suitable 'lag phase' to allow the transduced EC to integrate functionally in to the tumor vasculature. The two-step enzyme-prodrug 30 system offers greater flexibility of delicate clinical management, because cessation of GCV infusion in the event of (potentially very serious) complications arising from damage to normal EC, would block toxicity without the need to block transgene activity in situ.

A number of alternative 'suicide genes' in addition to thymidine kinase may also be useful for cancer gene therapy (Moolten et al., supra.). Introduction of the bacterial cytosine deaminase gene (Huber et al., 1993, Cancer Res. 53:4619) into tumor cells confers sensitivity to the antifungal agent 5-5 fluorocytosine (5-FC). Cytosine deaminase converts 5-FC to 5-fluorouracil (5-FC). FU, Nishiyama et al., 1985, Cancer Res. 45:1753). Since 5-FU is commonly used chemotherapeutic drug for breast cancer, several groups have developed cytosine deaminase-based 'suicide gene' therapy models for this disease. Tumor specificity may be further increased by introducing the c-erbB2 promoter/enhancer elements 5' to the cytosine deaminase gene, so that the 10 therapeutic transgene is preferentially transcribed in c-erbB2-overexpressing breast tumor cells (Harris et al., 1994, Gene Therap. 1:170). Alkaline phosphatase has been widely explored as prodrug-activating enzyme in the related field of antibody directed enzymne-prodrug therapy (ADEPT). This 15 enzyme has the advantage that it can activate a wide range of phosphorylated derivatives of anticancer agents (e.g. mitomycin C, etoposide, etc.) that cannot cross cell membranes until the charged phosphate group is cleaved off, so a single enzyme could generate de novo a cocktail of chemotherapeutic agents within the tumor mass (Senter et al., 1993, Bioconjugate Chem. 4:3). Other 20 suicide genes may encode a polypeptide or polypeptides (with a corresponding non-cytotoxic agent) such as Herpes Simpex virus thymidine kinase (gancyclovir or acyclovir), Varicella Zoster virus thymidine kinase (6 methoxypurine arabino nucleoside; Huber et al., 1991, Proc. Natl. Acad. Sci. USA 88:8039), E. coli cytosine deaminase (fluorouracil; Mullen et al., 1992, Proc. Natl. Acad. Sci. 25 <u>USA.</u> 89:33), E. coli xanthine-guanine phophoribosyl transferase (thioxanthine; Beshard et al., 1987, Mol. Cell Biol. 7:4139), E. coli or Leishmania purine nucleotide phosphorylase (various nontoxic purine deoxyadenosine, adenosine, deoxyguanosine, or guanosine derivatives (Koszalka and Krenitsky, 1979, J. Biol <u>Chem 254</u>:8185, 1979; Sorscher et al., 1994, <u>Gene Therapy 1</u>:233), cytochrome pla50 2B1 or cytochrome p450 reductase (e.g., 3amino-1,2,4 benzotriazine 1,4-30 dioxide (Walton et al., 1992, Biochem. Pharmacol. 44:251), cell surface alkaline phosphatase (e.g., etoposide monophosphate; Senter et al., 1988, Proc. Natl. Acad. Sci. USA 85:4842, 1988), nitroreductase (e.g., metronidazole or

nitrofurantoin; Hof et al., 1988, Immunitat und Infektion 16:220), N-deoxyribosy transferase (1-deazapurine; Betbeder et al., 1989, Nucleic Acids Res 17:4217), pyruvate ferrodoxin oxidoreductase (metronidazol; Upcroft et al., 1990, Int. J. Parasitolog, 20:489), carboxypepidase G2 (aminoacylate nitrogen mustards;

- Antoniw et al., 1990, Brit J. Cancer 62:909), carboxypeptidase A (methotrexate alpha alanine; Haenseler et al., 1992, Biochemistry 31:891), lactamase (cephalosporin derivatives; Meyer et al., 1993, Cancer Res. 53:3956; and Vradhula et al., 1993, Bioconjugate Chemistry 4:334), Actinomycin D synthetase complex (synthetic pentapeptide lactone precursors; Katz et al., 1990,
- J. Antibiotics 43:231), and •-glucuronidase (various glucuronide derivatives of toxic drugs such as doxorubicin; Bosslet et al., 1994, Cancer Res. 54:2151;
 Haeberlin et al., 1993, Pharmaceutical Res. 10:1553).

Any of a variety of other enzymes which convert inactive prodrugs into active drugs and known to those of skill in the art can also used in the gene delivery vehicles of the invention. For example, see PCT publication number WO 95/14014091 entitled "Compositions and Methods for Utilizing Conditionally Lethal Genes", and European Patent publication number EP90309430, entitled "Molecular Chimeras Useful for Cancer Therapy--Comprising Regulatory Sequences and heterologous enzyme, e.g. Varicella Zoster Virus Thymidine Kinase" for a description of additional prodrug/enzyme systems useful for gene therapy. As an additional example, see PCT Patent Publication No. WO 95/13095 entitled "New Prodrugs and Enzyme Targeting Molecule Conjugates--Useful in Antibody Direct Enzyme Prodrug Therapy of e.g. Viral Infections".

A variety of tumors may be targeted for treatment by the gene delivery vehicles of the invention. In general, solid tumors are preferred, although leukemias and lymphomas may also be treated if they have developed a solid mass, or if suitable tumor associated markers exist such that the tumor cells can be physically separated from nonpathogenic normal cells. Representative examples of suitable tumors include melanomas, colorectal carcinomas, lung carcinomas (including large cell, small cell, squamous and adeno-carcinomas), renal cell carcinomas and breast adeno-carcinomas. Gene delivery vehicles expressing thymidine kinase and other prodrug converting enzymes are also

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useful in the treatment of autoimmune diseases including rheumatoid arthritis, osteoarthritis and graft vs. host disease. See e.g. PCT Patent Publication No. WO 95/14091, entitled "Compositions and Methods for Utilizing Conditionally Lethal Genes," for a description of treatment of disease with gene therapy vectors expressing prodrug converting enzymes.

b. Cytokines

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A variety of polynucleotides encoding cytokines and immune system modulators can be delivered by the gene delivery vehicles of the invention for treatment of a number of different disorders. Representative examples include 10 cytokines, such as IL-1, IL-2 (Karupiah et al., 1990, J. Immunology 144:290-298; Weber et al., 1987, J. Exp. Med. 166:1716-1733; Gansbacher et al., 1990, J. Exp. Med. 172:1217-1224; U.S. Patent No. 4,738,927), IL-3, IL-4 (Tepper et al., 1989, Cell 57:503-512; Golumbek et al., 1991, Science 254:713-716, 1991; U.S. Patent No. 5,017,691), IL-5, IL-6 (Brakenhof et al., 1987, J. Immunol. 139:4116-4121; WO 90/06370), IL-7 (U.S. Patent No. 4,965,195), IL-8, IL-9, IL-10, IL-11, IL-12, IL-13 (Cytokine Bulletin, Summer 1994), IL-14 and IL-15, particularly IL-2, IL-4, IL-6, IL-12, and IL-13, alpha interferon (Finter et al., 1991, <u>Drugs</u> 42(5):749-765; U.S. Patent No. 4,892,743; U.S. Patent No. 20 4,966,843; WO 85/02862; Nagata et al., 1980, Nature 284:316-320; Familletti et al., 1981, Methods in Enz. 78:387-394; Twu et al., 1989, Proc. Natl. Acad. Sci. USA 86:2046-2050; Faktor et al., 1990, Oncogene 5:867-872), beta interferon (Seif et al., 1991, J. Virol. 65:664-671), gamma interferons (Radford et al., The American Society of Hepatology 2008-2015, 1991; Watanabe et al., 25 PNAS 86:9456-9460, 1989; Gansbacher et al., 1990, Cancer Research 50:7820-7825; Maio et al., 1989, Can. Immunol. Immunother. 30:34-42; U.S. Patent No. 4,762,791; U.S. Patent No. 4,727,138), G-CSF (U.S. Patent Nos. 4,999,291 and 4,810,643), GM-CSF (WO 85/04188), tumor necrosis factors (TNFs) (Jayaraman et al., 1990, J. Immunology 144:942-951), CD3 (Krissanen et al., 1987, 30 Immunogenetics 26:258-266, 1987), ICAM-1 (Altman et al., 1989, Nature 338:512-514; Simmons et al., 1988, Nature 331:624-627), ICAM-2, LFA-1, LFA-3 (Wallner et al., 1987, J. Exp. Med. 166(4):923-932), MHC class I molecules, MHC class II molecules, B7.1-.3, 2-microglobulin (Parnes et al.,

1981, <u>Proc. Natl. Acad. Sci. 78</u>:2253-2257), chaperones such as calnexin, MHC linked transporter proteins or analogs thereof (Powis <u>et al.</u>, 1991, <u>Nature</u> 354:528-531, 1991).

Genes encoding any of the cytokine and immunomodulatory proteins described herein can be expressed in a gene delivery vehicle of the invention. Other forms of these cytokines which are known to those of skill in the art can also be used. For instance, nucleic acid sequences encoding native IL-2 and gamma-interferon can be obtained as described in U.S. Patent Nos. 4,738,927 and 5,326,859, respectively, while useful muteins of these proteins can be obtained as described in U.S. Patent No.4,853,332. As an additional example, nucleic acid sequences encoding the short and long forms of mCSF can be obtained as described in US Patent Nos. 4, 847,201 and 4,879,227, respectively.

Other nucleic acid molecules that encode cytokines, as well as other nucleic acid molecules that are advantageous for use within the present invention, may be readily obtained from a variety of sources, including, for example, depositories such as the American Type Culture Collection (ATCC, Rockville, Maryland), or from commercial sources such as British Bio-Technology Limited (Cowley, Oxford England). Representative examples include BBG 12 (containing the GM-CSF gene coding for the mature protein of 127 amino acids), BBG 6 (which contains sequences encoding gamma interferon), ATCC No. 39656 (which contains sequences encoding TNF), ATCC No. 20663 (which contains sequences encoding alpha interferon), ATCC Nos. 31902, 31902 and 39517 (which contains sequences encoding beta interferon), ATCC No 67024 (which contains a sequence which encodes Interleukin-1b), ATCC Nos. 39405, 39452, 39516, 39626 and 39673 (which contains sequences encoding Interleukin-2), ATCC Nos. 59399, 59398, and 67326 (which contain sequences encoding Interleukin-3), ATCC No. 57592 (which contains sequences encoding Interleukin-4), ATCC Nos. 59394 and 59395 (which contain sequences encoding Interleukin-5), and ATCC No. 67153 (which contains sequences encoding Interleukin-6).

Gene delivery vehicles expressing the above cytokines are useful in the treatment of a variety of disorders. For example, see PCT publication number

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US94/02951 entitled "Compositions and Methods for Cancer Immunotherapy" for a description of gene therapy treatment of malignancy.

15. Neurological Disorders and Diseases

Polynucleotides encoding tyrosine hydroxylase can be useful in treating Parkinson disease.

For stroke or any acute brain injuries, polynucleotides encoding IGF-1, bFGF, vascular endothelial growth factor (VEGF) are useful.

16. Pulmonary Disorders

For treating emphysema, polynucleotides encoding α 1-anti-trypsin are useful.

For treating lung fibrosis, polynucleotides encoding superoxide dismutase (SOD) are useful.

For treating cystic fibrosis, polynucleotides encoding CFTR are useful.

15 ADDITIONAL AGENTS

Additional agents can be included with the desired polynucleotides to be delivered. These additional agents can facilitate endocytosis of the desired nucleic acids or aid binding of the nucleic acids to the cell surface or both, for example.

20 A. Polypeptides

One example are polypeptides which include, without limitation: asialoorosomucoid (ASOR); transferrin; asialoglycoproteins; antibodies; antibody fragments; ferritin; interleukins; interferons, granulocyte, macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), macrophage colony stimulating factor (M-CSF), stem cell factor and erythropoietin. Viral antigens, such as envelope proteins, can also be used. Also, proteins from other invasive organisms, such as the 17 amino acid peptide from the circumsporozoite protein of plasmodium falciparum known as RII.

B. Hormones, Vitamins, Etc.

Other groups that can be included are, for example: hormones, steroids, androgens, estrogens, thyroid hormone, or vitamins, folic acid.

C. Polyalkylenes, Polysaccharides, Etc.

Polyalkylene glycols can be included with the desired polynucleotides. In a preferred embodiment, the polyalkylene glycol is polyethlylene glycol. In addition, mono-, di-, or polysaccarides can be included. In a preferred embodiment of this aspect, the polysaccharide is dextran or DEAE-dextran. Also, chitosan and poly(lactide-co-glycolide)

D. <u>Lipids and Liposomes</u>

The desired polynucleotide can also be encapsulated in lipids or packaged in liposomes prior to delivery to the subject or to cells derived therefrom.

Lipid encapsulation is generally accomplished using liposomes which are able to stably bind or entrap and retain nucleic acid. The ratio of condensed polynucleotide to lipid preparation can vary but will generally be around 1:1 (mg DNA:micromoles lipid), or more of lipid. For a review of the use of liposomes as carriers for delivery of nucleic acids, see, Hug and Sleight, 1991, Biochim. Biophys. Acta. 1097:1-17; Straubinger et al., in METHODS OF ENZYMOLOGY (1983), Vol. 101, pp. 512-527.

Liposomal preparations for use in the instant invention include cationic (positively charged), anionic (negatively charged) and neutral preparations.

Cationic liposomes have been shown to mediate intracellular delivery of plasmid DNA (Felgner et al., 1987, Proc. Natl. Acad. Sci. USA 84:7413-7416); mRNA (Malone et al., 1989, Proc. Natl. Acad. Sci. USA 86:6077-6081); and purified transcription factors (Debs et al., 1990, J. Biol. Chem. 265:10189-10192), in functional form.

Cationic liposomes are readily available. For example, N[1-2,3-dioleyloxy)propyl]-N,N,N-triethylammonium (DOTMA) liposomes are available under the product line Lipofectin®, from GIBCO BRL, Grand Island, NY. (See, also, Felgner et al., 1987, Proc. Natl. Acad. Sci. USA 84:7413-7416). Other commercially available liposomes include transfectace (DDAB/DOPE) and

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DOTAP/DOPE (Boerhinger). Other cationic liposomes can be prepared from readily available materials using techniques well known in the art. See, e.g., Szoka et al., 1978, Proc. Natl. Acad. Sci. USA 75:4194-4198; PCT Publication No. WO 90/11092 for a description of the synthesis of DOTAP (1,2-

bis(oleoyloxy)-3-(trimethylammonio)propane) liposomes.

Similarly, anionic and neutral liposomes are readily available, such as from Avanti Polar Lipids (Birmingham, AL), or can be easily prepared using readily available materials. Such materials include phosphatidyl choline, cholesterol, phosphatidyl ethanolamine, dioleoylphosphatidyl choline (DOPC), dioleoylphosphatidyl glycerol (DOPG), dioleoylphoshatidyl ethanolamine (DOPE), among others. These materials can also be mixed with the DOTMA and DOTAP starting materials in appropriate ratios. Methods for making liposomes using these materials are well known in the art.

The liposomes can comprise multilamellar vesicles (MLVs), small
unilamellar vesicles (SUVs), or large unilamellar vesicles (LUVs). The various
liposome-nucleic acid complexes are prepared using methods known in the art.
See, e.g., Straubinger et al., in METHODS OF IMMUNOLOGY (1983), Vol.
101, pp. 512-527; Szoka et al., 1978, Proc. Natl. Acad. Sci. USA 75:4194-4198;
Papahadjopoulos et al., 1975, Biochim. Biophys. Acta 394:483; Wilson et al.,
1979, Cell 17:77; Deamer and Bangham, 1976, Biochim. Biophys. Acta
443:629; Ostro et al., 1977, Biochem. Biophys. Res. Commun. 76:836; Fraley et
al., 1979, Proc. Natl. Acad. Sci. USA 76:3348); Enoch and Strittmatter, 1979,
Proc. Natl. Acad. Sci. USA 76:145); Fraley et al., 1980, J. Biol. Chem.
255:10431; Szoka and Papahadjopoulos, 1978, Proc. Natl. Acad. Sci. USA
75:145; and Schaefer-Ridder et al., 1982, Science 215:166.

E. Lipoproteins

In addition, lipoproteins can be included with the polynucleotide to be delivered. Examples of lipoproteins to be utilized include: chylomicrons, HDL, IDL, LDL, and VLDL. Mutants, fragments, or fusions of these proteins can also be used. Also, modifications of naturally occurring lipoproteins can be used, such as acetylated LDL. These lipoproteins can target the delivery of polynucleotides to cells expressing lipoprotein receptors. Preferably, if

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lipoproteins are including with the polynucleotide to be delivered, no other targeting ligand is included in the composition.

If lipoproteins are included with the desired polynucleotides to be delivered, preferably, the composition comprises: (1) lipoprotein; (2) polynucleotide; and (3) a polynucleotide binding molecule.

Naturally occurring lipoproteins comprise a lipid and a protein portion. The protein portion are known as apoproteins. At the present, apoproteins A, B, C, D, and E have been isolated and identified. At least two of these contain several proteins, designated by Roman numerals, AI, AII, AIV; CI, CII, CIII.

A lipoprotein can comprise more than one apoprotein. For example, naturally occurring chylomicrons comprise A, B, C, and E, over time these lipoproteins lose A and acquire C and E apoproteins. VLDL comprises A, B, C, and E apoproteins, LDL comprises apoprotein B; and HDL comprises apoproteins A, C, and E.

The amino acids of these apoproteins are known and are described in, for example, Breslow, 1985, Annu Rev. Biochem 54:699; Law et al., 1986, Adv. Exp Med. Biol. 151:162; Chen et al., 1986, J Biol Chem 261: 12918; Kane et al., 1980, Proc Natl Acad Sci USA 77:2465; and Utermann et al., 1984, Hum Genet 65:232.

Lipoproteins contain a variety of lipids including, triglycerides, cholesterol (free and esters), and phopholipids. The composition of the lipids varies in naturally occurring lipoproteins. For example, chylomicrons comprise mainly triglycerides. A more detailed description of the lipid content of naturally occurring lipoproteins can be found, for example, in Meth. Enzym. 128 (1986). The composition of the lipids are chosen to aid in conformation of the apoprotein for receptor binding activity. The composition of lipids can also be chosen to facilitate hydrophobic interaction and association with the polynucleotide binding molecule.

Naturally occurring lipoproteins can be isolated from serum by ultracentrifugation, for instance. Such methods are described in Meth. Enzy., supra; Pitas et al., 1980, J. Biochem. 255:5454-5460; and Mahey et al., 1979, J. Clin. Invest 64:743-750.

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Lipoproteins can also be produced by *in vitro* or recombinant methods by expression of the apoprotein genes in a desired host cell. See, for example, Atkinson <u>et al.</u>, 1986, <u>Annu Rev Biophys Chem 15</u>:403, and Radding <u>et al.</u>, 1958, <u>Biochim. Biophys Acta</u> 30:443.

Lipoproteins can also be purchased from commercial suppliers, such as Biomedical Techniologies, Inc., Stoughton, Massachusetts, USA.

Mutants, fragments and fusion of the naturally occurring apoproteins are useful for delivery of polynucleotides. These polypeptides will retain more than about 80% amino acid identity; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polypeptides will exhibit more than about 92% amino acid sequence identity with naturally occurring lipoproteins or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99% sequence identity.

Such mutants, fragments and fusions can be constructed by altering the polynucleotides encoding the desired lipoproteins by recombinant DNA techniques. See, for example, Sambrook et al., (1989) Molecular Cloning, A Laboratory Manual, 2d edition (Cold Spring Harbor Press, Cold Spring Harbor, New York). These polynucleotides can be inserted into expression vectors and host cells can be utilized to produce the desired apoprotein.

In addition, naturally occurring lipoproteins, mutants, fragments, and fusions can be chemically altered. For example, acetylated LDL has biological activity. See, for example, Nagelkerke et al., 1983, J. Biol. Chem. 258(20):12221-12227; Weisgraber et al., 1978, J. Biol. Chem. 253:9053-9062; Voyta et al., 1984, J. Cell Biol. 99:2034-2040; Goldstein et al., 1979, Proc. Natl. Acad. Sci. USA 76:333-337; and Pitas, 1981, Arterosclerosis 1:177-185.

Chemically modified lipoproteins can also be purchased from commercial suppliers, such as Biomedical Techniologies, Inc., Stoughton, Massachusetts, USA.

All of these polypeptides exhibit receptor binding properties of naturally occurring lipoproteins. Usually, such polypeptides exhibit at least about 20% receptor binding of naturally occurring lipoproteins. More typically, the polypeptides exhibit at least about 40%, even more typically the polypeptides

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exhibit at least about 60%; even more typically, at least about 70%; even more typically, at least about 80%; even more typically, at least about 85%; even more typically, at least about 95% receptor binding of the naturally occurring lipoproteins.

Typically, lipoproteins are present in an amount effective to increase the frequency of incorporation of polynucleotides into a cell. Such an amount is sufficient to increase the frequency of incorporation of polynucleotides into a cell by at least 10%, compared to the frequency of incoporation of naked polynucleotides; more usually, at least 15%; even more usually, 20%; even more usually, at least 30%. The increase can be between 40 to 100%, and even 1000% and 10000% increase.

"Polynucleotide binding molecule" refers to those compounds that associate with polynucleotides, and the association is not sequence specific. For example, such molecules can (1) aid in neutralizing the electrical charge of polynucleotide, or (2) facilitate condensation of nucleotides, or (3) inhibit serum or nuclease degradation. Optionally, polynucleotide binding molecules can interact with lipoproteins by either hydrophobic association or by charge. Polynucleotide binding molecules include, without limitation, polypeptides, mineral compounds, vitamins, etc.

Examples of polynucleotide binding molecules include: polylysine, polyarginine, polyornithine, and protamine. Examples of organic polycations include: spermine, spermidine, and purtrescine. Other examples include histones, protamines, human serum albumin, DNA binding proteins, non-histone chromosomal proteins, coat proteins from DNA viruses, such as φX174, transcriptional factors also contain domains that bind DNA and therefore may be useful as nucleic aid condensing agents. Briefly, transcriptional factors such as C/CEBP, c-jun, c-fos, AP-1, AP-2, AP-3, CPF, Prot-1, Sp-1, Oct-1, Oct-2, CREP, and TFIID contain basic domains that bind DNA sequences.

Examples of other positively charged moieties include polybrene,
DEAE-dextran, and cationic lipids. Useful cationic lipids and liposomes are
described above. Lipids and liposomes are not used in this aspect of the
invention to encapsulate both polynucleotide and lipoprotein. The lipoprotein
must be exposed to bind the its cell surface receptor.

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Other synthetic compounds that are capable of binding negatively charged polynucleotides are useful, such as polymers of N-substituted glycines and others, as described below.

In a composition with a lipoprotein, the polynucleotide binding molecule can be present in an amount effective to neutralize the polynucleotide. However, the polynucleotide binding molecule also can be in excess of an effective amount to neutralize the polynucleotide to be delivered. Such an excess can produce a net positive electrical charge when complexed with the polynucleotides to be delivered. The positively charged complex can then interact with lipoproteins that comprise negatively charged lipids, such as phospholipids.

Typically, the polynucleotide binding molecule is in excess when the amount is 10% greater than the amount to neutralize the polynucleotide charge; more typically, the amount is 50% greater; even more typically, 100% greater; even more typically, 200% greater; even more typically, 200% greater; even more typically, 500% greater; even more typically, 20,000% greater; even more typically, 22,000% greater; even more typically, 25,000% greater; even more typically, 30,000% greater; even more typically, more than 40,000% greater than the amount effective to neutralize the electrical charge of the desired polynucleotide.

POLYCATIONIC AGENTS

Polycationic agents can be included, with or without lipoprotein, in a composition with the desired polynucleotide to be delivered.

Functional properties

A. Net Positive Charge

Polycationic agents typically exhibit a net positive charge at physiological relevant pH and are capable of neutralizing the electrical charge of nucleic acids to facilitate delivery to a desired location. These agents have both in vitro, ex vivo, and in vivo applications. For example, these polycationic agents can be used to transfect cells used to produce recombinant proteins.

Alternatively, the instant polycationic agents can be used to deliver nucleic acids to a living subject either intramuscularly, subcutaneously, etc.

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Physiological relevant pH varies somewhat between *in vitro* and *in vivo* applications. Typically, physiological pH is at least 5.5; more typically, at least 6.0; even more typically, at least 6.5. Usually, physiologically relevant pH is no more than 8.5; more usually, no more than 8.0; even more usually, no more than 7.5.

Preferably, the isoelectric point of the instant polycationic agents to neutralize nucleic acids is at least 9.

B. Non-Toxicity and Non-Immunogenic Properties

The composition of the polycationic agents of the invention will exhibit the toxicity and immunogenic properties desired. *In vitro* cell culture will have different immunogenic constraints than *in vivo* mammalian applications.

The instant polycationic agents can be easily tested for toxicity. For example, the agents can be added to medium for cells used in the *in vitro* assays, such as cos-7, Chinese Hamster Ovary cells, etc. Alternatively, the agents can be tested in standard animal tests for safety.

C. Condensation Properties

Due to the electric charge, a subset of these polycationic agents are capable of condensing the desired nucleic acids to a compact size to facilitate delivery. Typically, condensation "collapses" polynucleotides or nucleic acids into macromolecular structures, commonly into a toroid form. The smaller size of condensed nucleic acids eases delivery by facilitating, for example, packaging nucleic acids into liposomes and/or reducing exposure to proteases and/or nucleases.

The condensed nucleic acids exhibit different properties compared to "relaxed" nucleic acids, such as (1) a decrease in intercalation of ethidium bromide or other intercalating dye or (2) a reduced mobility in gel electrophoresis. Thus, condensation can be measured by at least two different assays, an intercalating dye assay or a band shift assay.

One type of intercalating dye assay uses ethidium bromide. In this assay, test nucleic acids, conveniently plasmid DNA, are mixed with polycationic agent in a ratio from about 1:1 to a 1:50 weight/weight ration of plasmid to condensing

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agent. Following incubation, ethidium bromide is added to the reaction to a final concentration of 1 µg/mL. If a nucleic acid such as RNA is used as the test nucleic acid, acridine orange may be used as the intercalating dye. The reaction mixtures are transferred into UV transparent plastic tubes spotted with 1% agarose gel, or placed upon UV transparent plastic c and illuminated with 260 nm light. The emission from the DNA-ethidium bromide complex is recorded on film by a camera equipped with an appropriate UV filter. The ability of an agent to condense DNA is inversely proportional to the intensity of the fluorescence in each reaction mixture.

The more precise test is a band shift assay. Briefly, this assay is performed by incubating nucleic acids, either labeled or unlabeled, with various concentrations of candidate condensing agents. Test nucleic acids, conveniently plasmid DNA, and condensing agent are mixed at 1:1 to 1:50 w/w ratios. Following incubation, each sample is loaded on a 1% agarose gel and electrophoresed. the gel is then either stained with ethidium bromide or dried and autoradiographed. DNA condensation is determined by the inability to enter the gel compared to a non-condensed standard. Sufficient condensation is achieved when at least 90% of the DNA fails to enter the gel to any significant degree.

Condensation can also be measure by directly determining the size of the complex using a light scattering instrument such as the a Coulter N4MD submicron analyzer, for example. Polynucleotides and a condensing agent are incubated at an appropriate ratio, either alone or in the present of 2% PEG-2000 (Fisher Scientific), and 0.6 M NaCl., and then diluted into 3 ml of water. This dilute solution is analyzed by the Coulter counter which will detect particles with a mean size of 0-1,000 nanometers (nm). Condensing agents, such as poly-Llysine, typically yield particles with a mean diameter of approximately 50-200 nm. See Lee et al., 1996, J. Biol. Chem. 271: 8481-8487.

D. Serum and/or Nuclease Protection Properties

The instant polycationic agents are capable of protecting nucleic acids from degradation in serum or from nucleases, including nucleases present in biological fluids, such as serum, prostate, synovial fluid, etc. One advantage of

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this type of protection is that smaller amounts of the desired nucleic acids are needed for efficient administration.

When present in effective amounts, these polycationic agents can inhibit serum degradation by at least 5 minutes as compared with uncomplexed nucleic acids; more usually, the amount used is sufficient to inhibit degradation by at least 10 minutes; even more usually; the amount used is sufficient to inhibit degradation by at least 30 minutes; even more usually, the amount used is sufficient to inhibit degradation by at least 45 minutes; even more usually, the amount used is sufficient to inhibit degradation by at least 60 minutes; even more usually, the amount used is sufficient to inhibit degradation by at least 90 minutes; and more usually, the amount used is sufficient to inhibit degradation by at least 120 minutes.

Increased serum protection can be measured simply by incubation of the polycation/polynucleotide complex with mouse serum, for example. Preferably, the serum will not be heat inactivated. After incubation, the mixture can be analyzed by gel electrophoresis to determine the quantity of the polynucleotides remaining after incubation.

Alternatively, nucleases can be added to the polycationic agent/nucleic acid complexes. The resulting mixture can be analyzed by gel electrophoresis to determine the amount of degradation. Other biological fluids, such as prostate flud, can also be tested.

E. Mediating Entry of Polynucleotides into a Cell

The polycationic agents can mediate entry of polynucleotides into a cell. Incorporation of polynucleotides into a cell can be measured by either protein expression assays or polynucleotide hybridization techniques, for example.

One method of detecting frequency of incorporation is to include a gene that encodes a marker protein, such as luciferase. Cells that have incorporated the delivered polynucleotides will express the marker protein. The protein can be detected by standard immunoassays, or by biological or enzymatic activity, as in the case of luciferase.

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Alternatively, standard hybridization techniques, such as Southern or Northern blots or polymerase chain reaction (PCR) techniques, can be used to detect the presence of the desired polynucleotides.

F. Additional Properties

- To facilitate entry of nucleic acids to the interior of cells, the instant agents can be capable of
 - (a) binding the polynucleotide to the cell surface;
 - (b) cell membrane destabilization;
 - (c) triggering endocytosis;
- 10 (d) endosome buffering capacity;
 - (e) releasing DNA/lipid complexes from endosomes; or
 - (f) nuclear tropism.

Assays for detecting these characteristics are standard and known to those skilled in the art.

15 Physical Properties

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The following physical characteristics are factors to consider when determining the composition of the polycationic agents:

- (a) distance between the substituents and the backbone
- (b) the total length of the chain;
- (b) hydrophobicity and/or aromacity;
 - (c) number of hydrogen bonding groups; and
 - (c) charge, including
 - (i) type of charge group, (ii) density of charge and (iii) position.

Other relevant characteristics include structural flexibility. For example, a

helical conformation of the polycationic agent may be preferred for some applications.

Specific dimensions to be considered include

- (a) the distance of phosphate groups in the polynucleotide of interest; and
- (b) the distance of monomer groups in the agents of interest.

Polypeptide Polycationic Agents

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The following are examples of useful polypeptides as polycationic agents: polylysine, polyarginine, polyornithine, and protamine. Other examples include histones, protamines, human serum albumin, DNA binding proteins, non-histone chromosomal proteins, coat proteins from DNA viruses, such as \$\phiX174\$, transcriptional factors also contain domains that bind DNA and therefore may be useful as nucleic aid condensing agents. Briefly, transcriptional factors such as C/CEBP, c-jun, c-fos, AP-1, AP-2, AP-3, CPF, Prot-1, Sp-1, Oct-1, Oct-2, CREP, and TFIID contain basic domains that bind DNA sequences.

Organic polycationic agents include: spermine, spermidine, and purtrescine.

The dimensions and of the physical properties of a polycationic agent can be extrapolated from the list above, to construct other polypeptide polycationic agents or to produce synthetic polycationic agents.

15 Synthetic Polycationic Agents

Synthetic polycationic agents which are useful include, for example,

DEAE-dextran, polybrene. Lipofectin®, and lipofectAMINE™ are monomers
that form polycationic complexes when combined with polynucleotides.

A preferred group of polycationic agents of the present invention have the following general formula (I):

$$Ta - \begin{bmatrix} R_1 & R_2 & O \\ | & V & | \\ -N - C - C - C - \end{bmatrix}_n - Tc$$

A preferred subset of these compounds include compounds having formula (I) where R₂ is hydrogen. Even more preferred are polymers comprising at least one natural amino acid. Also preferred are polymers where R₂ and R₃ are both hydrogen, also referred to as poly N-substituted glycines or poly NSGs.

A. Monomers

The polycationic agent of the invention comprises monomers with the following structure (Π):

$$\begin{array}{c|cccc}
R_1 & R_2 & O \\
 & V & || & & \\
 & -N - C - C - & & \\
 & R_3
\end{array}$$

Generally, R_1 , R_2 , and R_3 are moieties each with a molecular weight from 1 to 250 daltons. More typically, the molecular weight is no more than 200; even more typically, no more than 175.

Typically, each monomer comprises one hydrogen at R_1 , R_2 , or R_3 . More, typically, either R_1 and R_3 are both hydrogen, the structure of an L-amino acid; or R_2 and R_3 are both hydrogen, the structure of a NSG.

Monomers to be utilized in the polycationic agents can be either positively or negatively charged. Also, neutral substituents can also be utilized.

Degradation sites can be incorporated into the polymer, for example, by including substituents from a natural amino acid when R_1 and R_2 are hydrogen. These monomers can be positively or negatively charged, or neutral.

As a general rule, a basically charged monomer has a pKa value for the side chain of at least 7.5. Positively, or basically, charged monomers include without limitation those containing the following functional groups: amino, guanidino, hydrazido, and amidino. These functional groups can be either aromatic or aliphatic.

Positively charged monomers comprising hydrogen at R₃ and R₄, can be included in the polycationic agent, for example, as a degradation site. Such degradation site may aid in separation of the polycationic agent from the polynucleotide to permit further processing. For an L-amino acid like monomer, useful R₂ substitutents are, for example, from those found in naturally occurring amino acids, such as lysine and arginine. Also, sidechains from amino acid analogues can be used such as ornithine and canaline; or modifications of basic amino acids, such as homoarginine, and modifications of other amino acids such as guanidinovalinate, and aminoethylcysteine. The substitutents found in L-

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amino acids can also be incorporated at the R_1 and R_2 positions of the instant polycationic agents.

Naturally occurring amino acids and analogues are designated D-amino acids to indicate the chirality of these molecules. L-amino acids can also incorporated as monomers into the polycationic agents. The substituents of L-amino acids can be, for example, the same as those named for the D-amino acids.

Preferable monomers include N-substituted glycine monomers. Exemplary N-substitutions include alklphenyl, indolylalkyl, alkoxyphenyl, halophenylalkyl, hydroxyphenylalkyl, as well as the N-substitutions shown below.

$$\begin{array}{c|c} R_1 & R_3 \\ N-C-(CH_2)n-N-N+R_4 \\ R_2 & R_5 \end{array}$$

Alkylammonium,

where preferably $R_1 = H$;

 $R2 = H, CH_3;$

R3, R4, R5 can each be CH3, or CH3CH2; and n = 1-6.

$$\begin{array}{c|c} R_1 \\ N-C-(CH_2)n-N{H_3}^+ \\ R_2 \end{array}$$

Aminoalkyl,

where preferably $R_1 = H$;

 $R2 = H, CH_3;$

 $R_3 = CH_3$, CH_3CH_2 ; and

n = 1-6.

$$\begin{array}{c} R_1 \\ N-C \\ \downarrow \\ R_2 \end{array}$$

$$\begin{array}{c} NH_2 \\ NH_2 \end{array}$$

Guanidinoalkyl,

where preferably $R_1 = H$;

 $R2 = H, CH_3;$

 $R_3 = CH_3$, CH_3CH_2 ; and n = 1-6.

$$\begin{array}{c|c}
R_1 \\
N-C \\
-(CH_2)n \\
NH_2
\end{array}$$

$$\begin{array}{c|c}
NH_2^+ \\
NH_2
\end{array}$$

Amidinoalkyl,

where preferably $R_1 = H$;

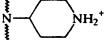
 $R2 = H, CH_3;$

 $R_3 = CH_3$, CH_3CH_2 ; and

n = 1-6.

$$N - NH_3^+$$

Aminocyclohexyl



Piperidyl

$$R_3$$
 R_4
 R_5
 R_5

Guanidinobenzyl

where preferably $R_1 = H$; and

 $R_2 = H, CH_3$

R3, R4, and R5 each can be H or __

$$R_1$$
 R_1
 R_2
 R_4
 R_5

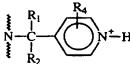
Amidinobenzyl

where preferably $R_1 = H$;

 $R_2 = H, CH_{3; and}$

R3, R4, R5 each can be -

NH₂



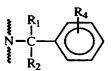
Pyridylmethyl

where preferably

 $R_1 = H$;

 $R_2 = H$, CH₃; and

 $R_4 = H, CH_3O, Cl, F, Br, CH, NO_2, CH_3.$



Aminobenzyl

where preferably

 $R_1 = H;$

 $R_2 = H$, CH₃; and

 $R_4 = H, CH_3O, Cl, F, Br, CH, NO_2, CH_3.$

The positively charged substituents described above can also be placed at the R₂ or R₃ positions of formulas (I) and (II).

The polycationic agents can comprise negatively charged or neutral monomers. As with the positively charged monomers, D-amino acid, L-amino acid, and NSGs are preferred to be incorporated as monomers.

The following are examples of such monomers:

B. Polycationic Polymers

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Typically, the polycationic agents exhibit a predicted isoelectric point of at least 9, excluding the terminal groups. Further, the agents contain, excluding the terminal groups, at least 20% positively charged monomers; more typically, at least 25% more typically, 30%; and preferably, at least 33% positively charged monomers. Typically, the agents do not comprises greater than 5% acidic monomers and preferably none.

The charge density and composition of the polycationic agent can be altered to accommodate the specific nucleic acid sequence, type, and other components included with the complex of nucleic acids and polycationic agent.

Usually, the length of the polymer is at least 8 monomers; even more usually, 12 monomers; even more usually, 18 monomers. More typically, the polycationic agents of the invention will be at least 24 monomer units in length; more typically, 30 monomer units; even more typically, 36 monomer units; even more typically, 48 monomer units. The polycationic agent can be up to 50 to 75 to 100 monomer units in length.

Preferably, the polycationic agent comprises monomers where all R_2 and R_3 are hydrogen. Even more preferably, where all R_2 and R_3 are hydrogen, the polycationic agent comprise repeating trimer units with the following monomer sequence (from amino to carboxy terminus): (1) neutral monomer, (2) neutral monomer, and (3) positively charged monomer.

Preferably, the neutral monomer comprises an aromatic group at the R₁ position; more preferably, wherein the aromatic group comprises a single ring; even more preferably, wherein the aromatic group is a six member ring.

Typically, the positively charged monomer is aminoalkyl at the R₁ position; more typically, the aminoalkyl comprises 1-6 carbon molecules; even more typically, the aminoalkyl is aminoethyl.

Typically, the polycationic agent comprises between 3 to 20 repeating trimers, trimers having two neutral and one positively charged R₁ groups are preferred, such as, for example, trimer shaving the sequence, neutral monomer, neutral monomer, positively charged monomer. More preferably, the polycationic agent comprises 5 to 18 trimers; preferably 8 to 16 trimers; and even more preferably, 12 to 16 trimers.

Optionally, the polycationic agent contains only positively charged monomers, excluding the terminal groups. Typically, such a polycationic agent comprises between 24 and 48 monomers; more typically, 30 to 40 monomers; even more typically, 36 monomers.

Polycationic agents of the present invention containing only positively charged monomers typically have guanidinoalkyl sidechains. Typically, the guanidinoalkyl sidechain comprises 1 to 6 carbon molecules. Preferably, the side chain is guanidino ethyl.

C. Neutral Polymers

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A preferred group of neutral polymers of the present invention have the general formula (I):

$$Ta - \begin{bmatrix} -R_1 & R_2 & O \\ -N - C - C - C - C \end{bmatrix}_n - Tc$$

Preferably, R₂ is hydrogen. Even more preferred are polymers comprising at least one natural amino acid. Also preferred are polymers having formula (I) where R₂ and R₃ are hydrogen, also referred to as poly N-substituted glycines or poly NSGs.

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Monomers employed in neutral polymers of the present invention have the same general formula as monomers employed in cationic polymers of the present invention, i.e.:

$$\begin{array}{c|cccc}
R_1 & R_2 & O \\
 & V & || \\
 & -N - C - C - C - C - C - R_3
\end{array}$$

Generally, R_1 , R_2 , and R_3 are moieties each with a molecular weight from 1 to 250 daltons. More typically, the molecular weight is no more than 200; even more typically, no more than 175.

Typically, each monomer comprises one hydrogen at R_1 , R_2 , or R_3 . More, typically, either R_1 and R_3 are both hydrogen, the structure of a L-amino acid; or R_2 and R_3 are both hydrogen, the structure of a NSG.

Monomers to be utilized in the neutral agents can be either positively or negatively charged. Also, neutral substituents can also be utilized. Neutral polymers exhibit no net positive or negative charge, excluding the terminal groups.

Degradation sites can be incorporated into the polymers by using naturally occurring amino acid substituents in monomers when R_1 and R_2 are hydrogen.

Naturally occurring amino acids and analogues are designated D-amino acids to indicate the chirality of these molecules. L-amino acids can also incorporated as monomers into the neutral polymers. The substituents of L-amino acids can be, for example, the same as those named for the D-amino acids.

Preferred monomers include N-substituted glycine monomers, and monomers that are capable of forming hydrogen bonds and/or ionic bonds with the polynucleotides to be delivered.

Examples of monomers for the neutral polymers include those described above and in the Examples below.

D. Linking Polymers Together

Polymers can be linked together incorporating terminating groups or sidechains that permit cross-linking of the polymers. For example, polymers can be linked by a disulfide bond. Other terminating groups useful for coupling polymers include, carbonate, urea, and the like.

E. Additional Groups to be Incorporated into the Polymer

Additional components can be included in the polycationic agents of the instant invention, such as targeting ligands. Such additional groups can facilitate endocytosis of the desired nucleic acids or aid binding of the nucleic acids to the cell surface.

Polypeptides can be incorporated into the polycationic agents. Examples include, without limitation: asialoorosomucoid (ASOR); transferrin; asialoglycoproteins; antibodies; antibody fragments; ferritin; interleukins; interferons, granulocyte, macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), macrophage colony stimulating

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factor (M-CSF), stem cell factor and erythropoietin. Viral antigens, such as envelope proteins, can also be used. Also, proteins from other invasive organisms are useful, such as the 17 amino acid peptide from the circumsporozoite protein of plasmodium falciparum known as RII.

In addition, lipoproteins can be incorporated into the polycationic agent, such as low density lipoprotein, high density lipoprotein, or very low density lipoprotein. Mutants, fragments, or fusions of these proteins can also be used.

Other groups that can be incorporated include without limitation: hormones, steroids, androgens, estrogens, thyroid hormone, or vitamins, folic acid. Folic acid can be incorporated into the polycationic agent according, for example, to Mislick et al., 1995, T.J. Bioconjugate Chem. 6:512.

Also, the polycationic agents of the instant invention can be chemically conjugated with polyalkylene glycol. In a preferred embodiment, the polyalkylene glycol is polyethlylene glycol. PEG can be incorporated with a polycation agent according, for example, to Lu et al., 1994, Int. J. Pept. Protein Res. 43:127.

In addition, the polycationic agent can be chemically conjugated with mono-, di-, or polysaccaride. In a preferred embodiment of this aspect, the polysaccharide is dextran.

These additional groups can be incorporated within the polycationic agent. For example, R_1 , R_2 , and R_3 can be a substituent that is capable of being activated to cross link with any one of the above groups. For example, a thiol group could be included to cross link with another group to form a disulfide bond.

F. Terminal Groups

The terminal groups of the instant polycationic agents can be chosen as convenient. Suitable terminal groups (i.e., Ta and Tc) include, for example, -NH₂, -OH, -SH, and -COOH. Terminal groups can be selected to enhance the targeting properties of the polycationic agent and can be any of the additional groups described above.

The additional groups described above can be incorporated at the terminus of the polycationic agent. For example, the polycationic agent can be

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(1) acylated with a variety of carboxylic acids; (2) sulfonylated with sulfonyl chlorides; or (3) derivatized with isocyanates or isothiocyanates. Once activated, the terminus can be reacted with any of the above-mentioned groups, such as a polypeptide, such as low density lipoprotein, or folic acid.

One means of adding a terminal group to the polycationic agent is, for example, is (1) to acylate the amino terminus with Fmoc-amino-hexanoic acid; and (2) to remove the protecting group, Fmoc, to generate a primary amine, which can be further functionalized.

Alternatively, the amino-terminal groups can include, without limitation: acyl, such as acetyl, benzoyl; or sulfonyl, such as dansyl.

Carboxy terminal groups can include, for example, amide or alkyl amide.

Synthesis of Polycationic Agents

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Polycationic agents of the present invention can be synthesized by either solid or solution phase methods. The following is a solid phase method for the synthesis of NSGs, which can be generally used for a wide variety of side-chain substitutents. This method can be performed utilizing automated peptide synthesis instrumentation to permit rapid synthesis of polycationic agents of interest. Such instruments are commercially available from, for example, Applied Biosystems and Milligen.

A. Two Step Monomer Assembly

A method of synthesis is to assemble the monomer from two submonomers in the course of extending a polymer comprising an NSG monomer. This technique is described in Zuckermann et al., 1992, J Amer Chem Soc 114(26):10646-10647, and Zuckermann et al., PCT Patent Publication No. WO 94/06451. The NSGs can also be considered to be an alternating condensation of copolymer of an acylating agent and an amine.

The direction of polymer synthesis with the submonomers occurs in the carboxy to amino direction. The solid-phase assembly for each monomer, in the course of polymer formation, eliminates the need for $N\alpha$ -protected monomers, as only reactive side-chain functionalities need to be protected.

Each monomer addition comprises two steps, an acylation step and a nucleophilic displacement step as shown in Figure 1.

Specifically, each cycle of monomer addition consists of two steps:

- (1) acylation of a secondary amine bound to the support with an acylating agent comprising a leaving group capable of nucleophilic displacement by an amine and a carbonyl group, preferably carboxyl. An example is a haloacetic acid; and
- (2) nucleophilic displacement of the leaving group with a sufficient amount of a submonomer comprising a primary amino group to introduce a sidechain. The amino group containing submonomer can be an alkoxyamine, semicarbazide, acyl hydrazide, substituted hydrazine or the like.

Acylation can be activitated with carbodiimide or other suitable carboxylate activation method.

The efficiency of the displacement is modulated by the choice of halide, e.g., I>Cl. Protection of aliphatic hydroxyl groups, carboxylic acids, carboxy, thiol, amino, some heterocycles, and other reactive side-chain functionalities is preferred to minimize undesired side reactions. However, the mild reactivity of some side-chain moieties toward displacement or acylation may allow their use without protection., e.g., indole, imidazole, and phenol.

B. Three Step Monomer Assembly

NSGs can also be constructed utilizing a three step method for assembling each monomer as the polymer is extended. The backbone of the monomer of first extended by acylation step followed by a nucleophilic displacement. The side chain is introduced by a second acylation step. The reaction scheme is shown in Figure 2.

The backbone of the monomer is assembled in the first two steps of the synthesis cycle. The first reaction is an acylation step where the carbonyl group of the acylating agent reacts with an amine. The acylating agent comprises a carbonyl group; a backbone, R_s; and a leaving group, L. Preferably, the carbonyl group is carboxyl.

The second step is a nucleophilic displacement of the leaving group by the first amino group of the displacing agent. The displacing agent comprises a

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first and a second amino group and a backbone, R_d. The first amino group is a primary amine, and the second step produces a secondary amine.

The third step is another acylation in which the another acylating submonomer reacts with the first amino group of the displacing agent to produce a tertiary amide. The acylation agent comprises a carbonyl group; an optional linker; and a sidechain. Preferably, the carbonyl group is carboxyl.

Pharmaceutical Compositions

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The polycationic agent/polynucleotide complexes, whether or not encapsulated in liposomes, may be administered in pharmaceutical compositions. The pharmaceutical compositions comprise a therapeutically effective amount of nucleic acids.

The term "therapeutically effective amount" as used herein refers to an amount of a therapeutic agent sufficient to detectably treat, ameliorate, or prevent a particular disease or condition, i.e., an amount sufficient to induce a detectable therapeutic or preventative effect. The effect may include, for example, chemical markers or antigen levels. Therapeutic effects also include reduction in physical symptoms, such as decreased body temperature. The precise effective amount for a subject will depend upon the subject's size and health, the nature and extent of the cardiovascular condition, and the therapeutics or combination of therapeutics selected for administration. Thus, it is not useful to specify an exact effective amount in advance. However, the effective amount for a given situation can be determined by routine experimentation and is within the judgment of the clinician. For purposes of the present invention, an effective dose will be from about 0.01 mg/kg to 50 mg/kg or 0.05 mg/kg to about 10 mg/kg of the DNA constructs in the individual to which it is administered.

A pharmaceutical composition can also contain a pharmaceutically acceptable carrier. The term "pharmaceutically acceptable carrier" refers to a carrier for administration of a therapeutic agent, such as antibodies or a polypeptide, genes, and other therapeutic agents. The term refers to any pharmaceutical carrier that does not itself induce the production of antibodies harmful to the individual receiving the composition, and which may be administered without undue toxicity. Suitable carriers may be large, slowly

metabolized macromolecules such as proteins, polysaccharides, polylactic acids, polyglycolic acids, polymeric amino acids, amino acid copolymers, and inactive virus particles. Such carriers are well known to those of ordinary skill in the art.

Pharmaceutically acceptable salts can be used therein, for example, mineral acid salts such as hydrochlorides, hydrobromides, phosphates, sulfates, and the like; and the salts of organic acids such as acetates, propionates, malonates, benzoates, and the like. A thorough discussion of pharmaceutically acceptable excipients is available in *Remington's Pharmaceutical Sciences* (Mack Pub. Co., N.J. 1991).

Pharmaceutically acceptable carriers in therapeutic compositions may contain liquids such as water, saline, glycerol and ethanol. Additionally, auxiliary substances, such as wetting or emulsifying agents, pH buffering substances, and the like, may be present in such vehicles. Typically, the therapeutic compositions are prepared as injectables, either as liquid solutions or suspensions; solid forms suitable for solution in, or suspension in, liquid vehicles, prior to injection may also be prepared. Liposomes are included within the definition of a pharmaceutically acceptable carrier.

Delivery Methods

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Once formulated, the compositions of the invention can be administered (1) directly to the subject; (2) delivered ex vivo, to cells derived from the subject; or (3) in vitro for expression of recombinant proteins. The subjects to be treated can be mammals or birds. Also, human subjects can be treated.

Direct delivery of the compositions will generally be accomplished by injection, either subcutaneously, intraperitoneally, intravenously or intramuscularly or delivered to the interstitial space of a tissue. The compositions can also be administered into a tumor or lesion. Other modes of administration include oral and pulmonary administration, suppositories, and transdermal applications, needles, and gene guns or hyposprays. Dosage treatment may be a single dose schedule or a multiple dose schedule.

Methods for the ex vivo delivery and reimplantation of transformed cells into a subject are known in the art and described in e.g., International Publication No. WO 93/14778 (published 5 August 1993). Examples of cells useful in ex

vivo applications include, for example, stem cells, particularly hematopoetic, lymph cells, macrophages, dendritic cells, or tumor cells.

Generally, delivery of nucleic acids for both ex vivo and in vitro applications can be accomplished by the following procedures, for example, dextran-mediated transfection, calcium phosphate precipitation, polybrene mediated transfection, protoplast fusion, electroporation, encapsulation of the polynucleotide(s) in liposomes, and direct microinjection of the DNA into nuclei, all well known in the art.

The examples presented below are provided as a further guide to the
practitioner of ordinary skill in the art, and are not to be construed as limiting the invention in any way.

Example 1

Synthesis of Polycationic Agents

This example describes the synthesis of polycationic agents with the following structure:

$$Ta - \left[-\frac{R_1}{N} - \frac{R_2}{C} - \frac{O}{C} - \frac{O}{C} - \frac{O}{C} \right]_n - Tc$$

where R₃ and R₂ are hydrogen for all monomers. All polymers describe in this example terminate in an amino and a carboxyl group unless specified, such as a folate terminating group.

The polycationic agents described below were synthesized according to the procedures described in Figliozzi et al., 1996, Meth. Enzy. 267:437-447, and Zuckermann et al., 1992, J. Amer. Chem. Soc. 114(26):10646-10647.

All polymers were synthesized using bromoacetic acid and primary amines. The following are substitutents of the primary amines to be positioned at R, to construct the polycationic agents:

Cationic Sidechains

Other Sidechains

$$H = (S)-\alpha$$
-methylbenzyl

H+ = (R)-
$$\alpha$$
-methylbenzyl

Nm = naphthylmethyl (R = H)
$$SN = (S) - \alpha - methylmaphthylmethyl (R = CH_3)$$

Other Sidechains

Peptoid-Folic acid conjugates

- Fmoc-aminohexanoic acid
 2... 20% piperidine/DMF

- 0.1 M Folic acid in DMSO, DIC HOBI, 50°
 Trifluoroacetic acid

Abbreviation	Description
Bn	benzyl
Chm	cyclohexylmethyl
Ff	furfurylmethyl
G	guanidinoethyl
Gp	guanidinopropyl
Н	(S) alpha-methylbenzyl
H+	(R) alpha-methylbenzyl
Ме	methoxyethyl
Nm	naphthylmethyl
P	aminoethyl
P'	aminohexyl
P*	(S)-α-methylaminoethyl
Ph	phenethyl
Pr	aminopropyl
Ру	N-pyrrolidinopropyl
Tmb	3,4,5,-trimethoxybenzyl
Q	trimethylaminoethyl
Phpr	phenylpropyl
6-gal	6-galactosyl
Тгр	N-2-(3-indolylethyl)
pMeOph	p-methoxyphenethyl
pClPh	p-chlorophenethyl
Tyr	p-hydroxyphenethyl
sN	(S)-α-methylnaphthylmethyl

The polycationic agents synthesized include:

Name	Sequence	Length	Mol. Wt.	# charges
RZ110-1	(HHP)6	18	2550.8	7
RZ110-2	(HP)9	18	2367.8	10
RZ110-3	(HPP)6	18	2184.8	13
RZ110-4	(HPPP)4HP	18	2123.8	14
RZ110-5	(HHP')6	18	2869.8	7
RZ110-6	(HP')9	18	2871.8	10
RZ110-7	(HP'P')6	18	2856.8	13
RZ110-8	(HP'P'P')4HP'	18	2851.8	14
RZ110-9	(HHP)12	36	5084.6	13
RZ110-10	(HP)18	36	4718.6	19
RZ110-11	(HPP)12	36	4352.6	25
RZ110-12	PP(HPPP)8HP	36	4230.6	27
RZ110-13	(HHP')12	36	5722.6	13
RZ110-14	(HP')18	36	5726.6	19
RZ110-15	(HP'P')12	36	5696.6	25

Name	Sequence	Length	Mol. Wt.	# charges
RZ110-16	P'P'(HP'P'P')8HP'	36	5686.6	27
RZ112-1	(Q)36	36	5181.5	37
RZ112-2	(G)36	36	5130.8	37
RZ112-3	(HP*P*P*)9	36	4529.3	28
RZ112-4	(P*)36	36	4122.8	37
RZ112-5	(HP*P*P*)4HP*	18	2305.2	14
RZ112-6	(P*)18	18	2069.9	19
RZ112-7	(P)18	18	1817.7	19
RZ112-8	(P)36	36	3618.4	37
RZ120-1	(MeMeP)8	24	2658.4	9
RZ120-2	(BnBnP)8	24	3170.8	9
RZ120-3	(HHP)8	24	3394.8	9
RZ120-4	(H+H+P)8	24	3394.8	9
RZ120-5	(MeMeP)12	36	3979.2	13
RZ120-6	(BnBnP)12	36	4747.6	13
RZ120-7	(HHP)12	36	5083.6	13
RZ120-8	(H+H+P)12	36	5083.6	13
RZ120-9	(MeMeP)16	48	5299.9	17
RZ120-10	(BnBnP)16	48	6324.5	17 .
RZ120-11	(HHP)16	48	6772.5	17
RZ120-12	(H+H+P)16	48	6772.5	17
RZ120-13	(HHP)12 folate	36	5300	13
RZ123-1	(HHPr)12	36	5252	13
RZ123-2	(HHPr)12	36	5252	13
RZ123-3	(HHP)12	36	5084	13
RZ123-4	folate-(HHPr)12	36	5862	13
RZ123-5	(HHGp)12	36	5756	13
RZ123-6	(HHG)12	36	5588	13
RZ124-1	(HHPr)12	36	5252	13
RZ124-2	(sNsNPr)12	36	6452	13
RZ124-3	(NmNmPr)12	36	6116	13
RZ124-4	(PyPyPr)	36	5756	13
RZ124-5	(HHPy)12	36	6069	13
RZ124-6	(Py)36	36	6573	1
RZ124-7	folate-(HHPr)12	36	5862	13
RZ127-1	(PhPhP)12	36	5085	13
RZ127-2	(ChmChmP)12	36	4895	13
RZ127-3	(TmbTmbP)12	36	6912	13
RZ127-4	(FfFfP)12	36	4508	13
RZ136-3	(PhprPhprP)12	36	5419	13
RZ140-2	(6-gal)12-(PhPhP)12	48	7712	13
RZ140-3	(TrpTrpP)12	36	6020	13
RZ144-1	(PhPPh)12	36	5083	13
RZ144-2	(PPhPh)12	36	5083	13
RZ144-3	(pMeoPhpMeoPhP)	36	5803	13

Name	Sequence	Length	Mol. Wt.	# charges
RZ144-4	(pClPhpClPhP)12	36	5910	13
RZ144-5	AMCA-(PhPhP)12	36	5411	12
RZ144-8	(TyrTyrP)12	36	5467	12
RZ144-12	(6gal6galP)12	36	6475	13
*RZ145-1	(PhPhP)12	36	5085	13
RZ147-2	(PpMeOPhpMeOPh)12	36	5805	13

^{*}purified

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To summarize the method, Fmoc-Rink amide resin (NovaBiochem, San Diego, California, USA) is used as the solid support. This is the same resin that is used for the Fmoc synthesis of peptide C-terminal amides. The polycationic synthesis begins with the deprotection of the Fmoc group on the resin with 20% (v/v) piperidine-dimethylformamide (DMF). The amino resin is then acylated with bromoacetic acid. This is followed by nucleophilic displacement of the bromide with a primary amine to build the NSG monomer. The latter two steps are then continued in an iterative fashion to elaborate the desired oligomer.

All reactions and washings were performed at room temperature unless otherwise noted. Washing of the resin refers to the addition of a wash solvent (usually DMF or dimethylsulfoxide (DMSO)) to the resin, agitating the resin so that a uniform slurry is obtained (typically for about 20 seconds), followed by thorough draining of the solvent from the resin. Solvents were removed by vacuum filtration through the fritted bottom of the reaction vessel until the resin appeared dry (typically about 5 seconds). In all the syntheses, resin slurries were agitated via bubbling argon up through the bottom of the fritted vessel.

A fritted reaction vessel was charged with 100 mg (50 μ mol) of Fmoc-Rink amide resin with a substitution level ~ 0.50 mml/g resin. Two milliliters of DMF was added to the resin and this solution was agitated for 1-2 minutes to swell the resin. The DMF was then drained. The Fmoc group was then removed by adding 2.0 ml of 20% piperidine in DMF to the resin. This was agitated for 1 minute and then drained. Another 2 ml of 20% piperidine in DMF was added to the resin and agitated for 15 minutes and then drained. The resin was then washed with DMF, six times with 2 ml.

The deblocked amine was then acylated by adding 850µl of 0.6 M bromoacetic acid in DMF to the resin followed by 200 µl of 3.2 M N,N'-

diisoprooplycarbodiimide (DIC) in DMF. This solution was agitated for 30 minutes at room temperature and then drained. This step was repeated a second time. The resin was then washed with DMF, twice with 2 ml and DMSO, once with 2 ml. This completed one reaction cycle.

The second cycle was initiated by the acylating step with bromoacetic acid and DIC, followed by displacement with the second amine. This acylation/displacement cycle was repeated until the desired oligomer was obtained.

Cleavage of the resin from the polycationic agent is as follows. The dried resin was placed in a glass scintillation vial containing a teflon-coated micro stir bar, and approximately 5 ml of 95% trifluoroacetic acid (TFA) in water was added. The solution was stirred for 20 minutes and then filtered through an 8-ml solid-phase extraction (SPE) column fitted with a 20-µm polyethylene frit into a 50 ml polypropylene conical centrifuge tube.

The resin was washed with 1 ml 95% TFA. The combined filtrates were then lyophilized three times from 1:1 acetonitrile:water. Material was redissolved to a concentration of 5 mM in 5% acetonitrile in water.

Preparation of guanidinoalkyl-containing Polymers:

The guanidinoalkyl sidechains were introduced into the polymers by post-synthesis modification of aminoalkyl sidechains. Thus, polymers were synthesized by the submonomer method as described above except that methoxybenzhydrylamine (MBHA) resin was used instead of the Rink resin. Wherever a guanidinoalky sidechain was desired, a mono-Boc-alkanediamine was incorporated in the displacement step. After elaboration of the polymers, the sidechain Boc groups were removed by treatment with 95% TFA/water for 20 min at room temp. (This does not remove the oligomer from the soliud support). The free amino groups were then guanidinylated by treatment with 1 H-pyrazole-1-carboxamidine (1 M in DMF, 2 x 1 hr, 40°C). After washing with DMF and methylene chloride, the oligomer was cleaved from the resin with hydrofluoric acid, and lyophilized.

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Preparation of folic acid - polymer conjugates:

Folic acid - polymer conjugates were prepared by adding a linker to the N-terminus of the resin-bound polymer which was then acylated with folic acid. Specifically, after elaboration of the polymer, the N-terminus was acylated with Fmoc-aminohexanoic acid (0.5 M in DMF, 0.5 M hydroxybenzotriazole, 0.5 M diisopropylcarbodiimide (DIC), 1 x 1 hr, room temp.). After Fmoc group removal (20% piperidine/DMF, 1 x 20 min, room temp.), the free primary amino group was acylated with folic acid (0.1 M in DMSO, 0.1 M DIC, 1 x 2 hr, 50°C). After washing of the resin, the conjugate was cleaved with 95% TFA/water in the usual fashion.

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Example 2

Condensation of Polynucleotides

Polycationic agents were synthesized and isolated to a final concentration of 5 mM as described in Example 1. Polynucleotides were condensed with RZ110, RZ112, and RZ120 series compounds according to the following procedure.

- (1) Dilute all polycationic agents to a final concentration of 3 nanomoles of positive charge per microliter.
 - (2) Add 1 µg of DNA to 1-2µl of diluted polycationic agents.
- (3) Adjust volume to 10 μ l. This mixture can be stored overnight at 20 4°C.
 - (4) Add of 5 μ l of DNA/polycationic mixture to 2 μ l of 5X buffer, which does not contain SDS to maintain the complex. (5X buffer = 40% sucrose, 0.25% bromphenol blue and 200 mM Tris Acetate, 4 mM EDTA (PH 7.8).
 - (5) Adjust volume to $10 \mu l$.
 - (6) Run sample on a 1% agarose gel utilizing 75 volts for 1.5 hours...

Between 1 to 2 μ l, all polycationic agents were judged to retard the migration of DNA into an agarose gel.

Example 3

Inhibition of Serum Degradation

The RZ110, RZ112, and RZ120 series compounds were mixed with polynucleotide as described in Example 2. Five microliters of the overnight mixture was added to 5 µl of BalbC mouse serum. The serum was not heat treated but freeze thawed. The serum, polycationic agent, and polynucleotide mixture was incubated typically for 30 minutes at 37°C. The time of incubation varied between 5 and 60 minutes

Next, 2 μ l of 5X buffer containing 0.5% (wt/v) SDS was added to the incubated mixture. This final solution was loaded onto a 1% agarose gel and electrophoresed at 75 volts for 1.5 hours.

All of the compounds tested, i.e., the entire RZ110, 112, and 120 series, provided significant protection in a direct comparison. The entire RZ112 series and RZ110-3 and RZ110-8 inhibited serum degradation better than poly-Llysine.

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Example 4

Peptoid Mediated In Vitro Delivery

DNA comprising a luciferase gene $1\mu g/\mu l$, was diluted into endotoxin free water. The plasmid DNA was CMVKm luciferase, which is described in more detail in Example 5.

The transfection protocol for in vitro delivery was as follows:

- (A) HT1080 cells were used. These cells are available from American Type Culture Collection, Rockville, Maryland, USA, Accession No. CCL 121. This is a fibrosarcoma. The growth medium was Dulbecco's Modified Eagle medium (DME) with 10% heat-inactivated fetal calf serum.
- 25 (B) Twenty four hours prior to transfection, the cells were placed at 5 X 10⁴ per well of a 24-well plate in 1 ml of medium.
 - Feed cells with 500μl of DME-10% fetal calf serum (FCS)or 500μl
 Opti-MEM®. Opti-Mem® can be purchased from Gibco BRL, Life
 Technologies, Inc., Gaithersburg, Maryland, USA.
- 30 2. Add 200 μl Opti-MEM® to each tube.

- 3. Add 3 µl of the desired polycationic agent to the 200µl of Opti-MEM®.
 - Add 2 µl of 1µg/µl luciferase DNA, mix.
 - 5. Incubate mixture for 5 minutes at room temperature.
- 5 6. Add 100 µl of the polycationic agent/DNA mixture to plate with DME-FCS, 100 µl to cells fed with Opti-MEM®.
 - 7. Incubate cells and polycationic agent/DNA mixture for ~4 hours at 37°C.
 - 8. Change media on all cells to DME-FCS.
 - DME-FCS was used as a positive control. As a control, a transfectant, LT1, was used from Panvera, Inc., Madison,

Wisconsin, USA to transfect cells in serum and cells in Opti-MEM®.

Cells were tested for luciferase activity using a Promega Luciferase Assay System from Promega, Madison, Wisconsin, USA., in accordance with 15 the manufacturer's directions.

Results:

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Name	Formula	Luciferase (RLU)
RZ120-1	(MeMeP)8	0
RZ120-2	(BnBnP)8	0.93
RZ120-3	(HHP)8	1.38
RZ120-4	(H+H+P)8	1.5
RZ120-5	(MeMeP)12	0
RZ120-6	(BnBnP)12	1.64
RZ120-7	(HHP)12	2.64
RZ120-8	(H+H+P)12	2.84
RZ120-9	(MeMeP)16	0
RZ120-10	(BnBnP)16	1.42
RZ120-11	(HHP)16	1.94
RZ120-12	(H+H+P)16	1.32
control	LT1	51.96

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Experiment #2		
Name	Formula	Luciferase
		(RLU)
RZ110-1	(HHP)6	0.0015
RZ110-2	(HP)9	0.0012
RZ110-4	(HPPP)4HP	0.0004
RZ110-5	(HHP')6	0.0006
RZ110-6	(HP')9	0.0052
RZ110-7	(HP'P')6	0.0005
RZ110-8	(HP'P'P')4HP'	0.0003
RZ110-9	(HHP)12	8.7
RZ110-10	(HP)18	0.0014
RZ110-12	PP(HPPP)8HP	0.0459
RZ110-13	(HHP')12	2.5
RZ110-14	(HP')18	2.2
RZ110-15	(HP'P')12	0.064
RZ110-16	P'P'(HP'P'P')8	0.01
control	LTI	88.7

Example 5

Targeting Ligand

A. Cells, Vector, and Compositions Used.

In a first experiment, murine endothelial cells (Py-4-1) which express high levels of acetylated-LDL receptors. The cells and the LDL receptors are described in Dubois et al., 1991, Exp. Cell Res. 196:302-313.

A luciferase-containing plasmid (pCMVkmLUC) was used to determine if polynucleotides could be delivered and expressed into endothelial cells when associated with polycationic agents described in Example 1 with acetylated-LDL (Ac-LDL). A description of the identification and isolation of endothelial cells based on their increased uptake of acetylated-low density lipoprotein is in Voyta et al., 1984, J. Cell Biol. 99: 2034-2040.

The plasmid used in these experiments pCMVkmLUC, was constructed by inserting the luc + gene from pSP-luc+ (Promega Corporation, Madison, WI) into the expression vector pCMVkm2. Briefly, pSP-luc+ was digested with the restriction enzymes Nhe1-EcoRV (Boehringer Mannheim, Indianapolis, IN) and a fragment of 1691 bp was isolated by standard methods. This fragment was inserted into pCMVkm2, which had been digested with XbaI and EcoRV using the Rapid Ligation Kit (Boehringer Mannheim, Indianapolis, IN). The sequence of pCMVkm2 is depicted in SEQ ID NO:2 and described below. The luc + gene was cloned into pCMVkm2 such that expression is driven by the CMV immediate early enhancer promoter and terminated by the bovine growth hormone polyadenylation signal.

The luciferase expression was compared to levels obtained with the same vector delivered in conjunction with lipofectamine, an agent used commonly to transfect cells *in vitro* (Hawley-Nelson <u>et al.</u>, 1993, <u>Focus 15:73</u>). The results are presented in the table below.

B. Method of Transfection:

Briefly, the cells were plated in 24 well dishes, grown to approximately 80% confluence, transfected and assayed 24 hours later for luciferase activity. All transfections were done in serum containing medium. During transfection mixture preparation, pCMVkmLUC was first mixed with RZ 112, and the DNA-

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cationic polycationic agent complexes were then added to Ac-LDL. Serum containing medium was then added to the mixtures to adjust the volume delivered to each well to 0.5 ml.

Lipofectamine was used as a positive control. No lipoprotein was added to this positive control. Lipofectamine is a 3:1 (w/w) liposome formulation of the polycationic lipid 2,3,-dioleylosy-N-[2(spermine-carboxamido)ethyl]-N,N-dimethyl-1-propanaminiumtrifluoroacetate (DOSPA) and the neutral lipid dioleoyl phosphatidyl-ethanolamine (DOPE) in membrane-filtered water. Lipofectamine can be purchased from Life Technologies, Gaithersburg, Maryland, USA).

C. Luciferase Assay

Luciferase activity was assayed using the Promega Luciferase Assay System, Madison, Wisconsin.

15 D. Results

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Table 1 shows the results of an experiment where the polycationic agent, RZ-112-2 was compared to lipofectamine to deliver the luciferase gene to cells comprising the acetylated LDL receptor.

Table 1
LuciferaseActivity

Group	Ac-LDL (μg)	pCMVkmLUC (µg)	RZ 112-2 (nm)	pg luc/mg protein
1	•	10	2.5	61
1	J	10	2.5	61
2	-	10	-	0
3	-	10	2.5	16
4	0.5	1	0.25	16
5	-	1	0.25	17
6	0.5	10	2.5	631
7	0.5	10	5	1996

LIPOFECTAMINE CONTROL

8 - 10 - 10786

(GC)

^{*} each number represents the mean of three wells.

Example 6

Comparison of Cells with and without Acetylated LDL Receptors

A. Cells with Acetylated LDL Receptors

For this experiment, K1735 mouse, epithelial melanoma cells were used.

These cells express low or non-existent levels of Ac-LDL receptors. A description of the cells is in <u>J. Natl. Cancer Inst.</u> 69(4): (1982).

B. Methods

Briefly, the cells were plated in 24 well dishes at 10,000 cells per well in DME with 10% FCS supplemented with 2 mM L-glutamine. The Py-4-1 cells were cultured in 10% CO₂ at 37°C. The K1735 cells were cultured in 5% CO₂ at 37°C. The cells were grown to approximately 50% confluence, transfected and assayed 24 hours later for luciferase activity. All transfections were done in serum containing medium.

During transfection mixture preparation, pCMVkmLUC was first mixed with RZ 112-2, and the DNA-polycationic agent complexes were then added to Ac-LDL. Serum containing medium was then added to the mixtures to adjust the volume delivered to each well to 0.5 ml.

C. Results

<u>Table 2</u> <u>Luciferase Activity</u>

Group	Ac-LDL (μg)	pCMVkmLUC (µg)	RZ 112-2 (nm)	pg luc/mg protein*	
				Py-4-1	K1735
1	0.5	1	1	1301	24
2	0.5	1	5	2181	0
3	0.5	1	10	373	0
4	0.5	10	5	840	0
5	-	1	5	327	0
6	-	1	5	945	ND
10	5	1	5	298	ND
LIPOFE	CTAMINE (CONTROL			
7	-	1	-	23	0
8	-	10	-	2878	960

^{*} each number represents the mean of three wells.

Example 7 Injection of Polynucleotides Encoding Erythropoietin

A. Polynucleotides

CMVkm2 is the standard vector used in these studies. CMVkm2 is a vector optimized for expression in mammalian cells. The gene of interest is cloned into a polylinker which is inserted 3' of a human CMV expression cassette. This cassette contains the human CMV immediate early promoter/enhancer followed by intron A of the human CMV immediate early region (Chapman et al., 1991, Nucl. Acids Res. 19:3937-3986). Transcription is terminated by a polyadenylation site from the bovine growth hormone gene, which has been cloned immediately 3' of the polylinker. See SEQ ID NO:2 for the CMVkm2 vector.

The CMV-km-cmEPO vector was constructed from CMVkm2 as follows. The cynomolgus monkey EPO cDNA was acquired from the ATCC (Accession No. 67545, Rockville, MD). This plasmid was cut with <u>AvrII</u> and <u>BglII</u> and inserted into the <u>XbaI</u> and <u>BamHI</u> sites of the CMVkm2 vector. The inserted sequence contains the entire coding region of cmEPO (Genbank accession M18189). See SEQ ID NO:3.

B. Mice

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Immunodeficient severe combined immunodeficiency (SCID) mice were obtained from Charles River Labs, Wilmington, Massachusetts, USA.

Intramuscular injections were performed as follows: mice were anaesthetized with 50 µl of a solution which contained 80 mg/ml ketamine and 4 mg/ml of xylazine. The area surrounding the anterior tibialis muscle was shaved. Fifty µl of DNA, at a concentration of 2.7 ug/µl in .9% saline solution was injected into the anterior tibialis muscle of both legs using a 28 gauge needle. Twenty-four hours after the first injection, a second injection was performed using the identical protocol. Blood was taken from the orbital sinus to determine hematocrits on a weekly basis.

C. Result

The hematocrit readings on 6 mice which were injected with plasmid, CMVkm-cmEpo (which expresses the cynomolgus monkeys EPO cDNA), are shown in Table 3 below. The row marked control shows the average reading for three uninjected mice. The raw data for the three control mice is shown in the lower part of Table 3. Mouse 2 in the injected group died between 4 and 5 weeks post-injection.

Table 3
Hematocrit Levels (%)

	Week 0	week 1	week 2	week 3	week 4	week 5	week 6	week 7	week8	week 9	week 10
Week#											
ſì											
Mouse ⊍											
mouse 1	05	63	5.99	57.5	63	63.5	56.5	62.5	54	53.5	54.5
mouse 2	20	64	29	56.5	55.5						
mouse 3	50	09	61.5	63	19	56	49.5	53	53.5	54.5	57
mouse 4	20	62	68.5	71.5	67.5	09	62.5	59.5	57.5	53.5	55.5
mouse 5	-20	62	62.5	99	19	53.5	58	52.5	54	52.5	48.5
mouse 6	20	99	63.5	62.5	09	58	58	53.5	55.5	52.5	52.5
Control	20	51.5	48	47.5	53	49.5	49.5	50.5	51.5	51	49
control	week 0	control									
		wk 1	wk 2	wk 3	week 4	week 5	week 6	week7	week 8	week 9	week 10
mouse 1	20	52	48	46.5	52.5	48	49.5	52	51.5	50.5	46.5
mouse 2	20	15	47	14	55	51	50.5	50	52	49	51.5
mouse 3	0\$	52	49	48.5	52	49.5	48.5	48.5	51	52.5	50

Example 8

Injection of Polynucleotides Encoding Leptin

A. Polynucleotides

The CMV-km2 vector, described above, was used for these experiments. Either the wild-type or HA version of the leptin coding region was inserted into the vector. The map of the plasmid is depicted in Figure 4 and the sequence of the vector with the wild type leptin is shown in SEQ ID NO:4.

B. Mice

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Ob/ob mice were obtained from Jackson Labs, Bar Harbor, Maine, USA. The first of the recessive obesity mutations, the *obese* mutation (*ob*) was identified and described in 1950 by Ingall et al., 1950, J. Hered. 41:317-318. Subsequently, 5 single-gene mutations in mice have been observed to produce an obese phenotype, as described in Friedman et al., 1990, Cell 69:217-220. (More recently, the mouse *obese* gene and its human homologue have been cloned, as described in Zhang et al., 1994, Nature 372:425).

C. Method

Intramuscular injections were performed as follows: mice were anaesthetized with the same ketamine solution described above in the Example 7 and the area surrounding the anterior tibialis muscle was shaved.

Fifty microliters of DNA at a concentration of 3.3. μ g/ μ l in 0.9% saline solution was injected into the anterior tibialis muscle of both legs using a 28 gauge needle.

Seventy-two hours after the first injection, a second injection was performed using the identical protocol.

Group 1 ob/ob mice were injected with a plasmid (CMVkM leptin-wt) which encodes the wild-type mouse leptin protein.

Group 2 ob/ob mice were injected with a plasmid (CMVkm-leptinHA) which encodes a form of mouse leptin which is modified with the epitope which is recognized by the antibody 12CA5. The amino acid sequence of the epitope is SYPYDVPDYASLGGPS (Wilson et al., 1984, Cell 37: 767-778).

Group 3 ob/ob mice were injected with a solution of 0.9% saline.

The mice were weighed each day (see Table 4) and the proportional weight gain for each mouse during the first eight days was calculated. The results are shown in Table 5. For any given day, the weight was subtracted from the weight of the individual mouse on day 0, and the difference was divided by the weight on day 0. The proportional weight change data from day 8 was analyzed using an unpaired t-test. When compared with group 3 control mice the p value from group 2 mice was 0.004. When compared with group 3 control mice, the p value for group 1 mice is 0.0038.

Note: the mice were not weighed on day 1 and day 2, the values for these days were extrapolated from day 3.

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Table 4

Weight of Mice in Grams	f Mice ir.	Grams									
group 1	day 0	day3	day4	day5	day6	day7	day8	dav9	dav 10	dav11	day 12
mouse 1	47	49	49	51	51	51	51	52	52	23	53
mouse 2	48	49	49	51	51	51	51	52	53	3 5	53
monse 3	46	48	48	49	49	49	49	50	25	3 5	21
mouse 4	47	48	48	49	49	49	50	20	20	S	51
mouse 5	49	50	50	51	51	51	52	52	52	52	52
group 2	day 0	day 3	day 4	day 5	day 6	day 7	day 8	day 9	day 10		
mouse 1	49	50	50	52	52	52	52	53	22	24	3,5
mouse 1	43	45	44	45	45	4	45	45	45	46	47
mouse 3	48	49	49	50	50	50	51	52	52	52	52
mouse 4	49	50	50	52	51	51	52	52	52	52	53
mouse 5	46	48	49	49	49	50	50	51	51	51	51
group 3	day 0	day 3	day 4	day 5	day 6	day 7	day 8	day 9	dav 10		
mouse 1	40	42	42	43	43	45	45	45		45	46
mouse 2	48	49	50	50	51	52	52	52	52	53	53
mouse 3	48	50	50	52	52	53	53	55	55	55	55
mouse 4	49	52	52	53	53	54	55	54	54	54	55
mouse 5	43	45	46	47	48	49	20	49	49	48	49

Table 5

Proportion (gp1), Le				n Day 0	of Mice	Injected	with cD	NA for L	eptin
Group 1 N	Mice Bel	ow Injec	ted with	CMVkM	1-Leptin	-wt			
	day 0	day I	day 2	day 3	day 4	day 5	day 6	day 7	day 8
mouse 1	0	0.009	0.018	0.028	0.028	0.059	0.059	0.059	0.066
mouse 2	0	0.013	0.026	0.039	0.039	0.085	0.085	0.085	0.085
mouse 3	0	0.006	0.012	0.018	0.018	0.062	0.062	0.062	0.062
mouse 4	0	0.014	0.028	0.042	0.042	0.065	0.065	0.065	0.065
mouse 5	0	0.007	0.014	0.021	0.021	0.042	0.042	0.042	0.06
	0	0.007	0.014	0.021	0.021	0.041	0.041	0.041	0.061
Group 2 N	Aice Bel	ow Injec	ted with	CMVkN	1-Leptin	HA			
	day 0	day 1	day 2	day 3	day 4	day 5	day 6	day 7	day 8
mouse 1	0	0.006	0.013	0.02	0.02	0.06	0.06	0.06	0.06
mouse 2	0	0.013	0.031	0.046	0.023	0.046	0.046	0.046	0.046
mouse 3	0	0.007	0.014	0.021	0.021	0.04	0.04	0.04	0.062
mouse 4	0	0.006	0.012	0.02	0.02	0.06	0.04	0.04	0.06
mouse 5	0	0.014	0.028	0.043	0.065	0.065	0.065	0.08	0.08
Group 3 N	Mice Bel	ow Injec	ted with	Saline					
	day 0	day l	day 2	day 3	day 4	day 5	day 6	day 7	day 8
mouse 1	0	0.016	0.032	0.05	0.05	0.06	0.06	0.125	0.125
mouse 2	0	0.07	0.014	0.021	0.04	0.04	0.06	0.08	0.08
mouse 3	0	0.013	0.026	0.04	0.04	0.08	0.08	0.1	0.1
mouse 4	0	0.02	0.04	0.06	0.06	0.08	0.08	0.1	0.12
mouse 5	0	0.015	0.03	0.045	0.069	0.09	0.12	0.14	0.14
Average I	roportio	nal chan	ge in We				,		
	day 0	day 1	day 2	day 3	day 4	day 5	day 6	day 7	day 8
group 1	0	0.009	0.018	0.028	0.028	0.059	0.059	0.059	0.066
group 2	0	0.009	0.018	0.028	0.029	0.054	0.05	0.053	0.061
group 3	0	0.014	0.028	0.042	0.051	0.07	0.08	0.109	0.113

Example 9

Peptoid Mediated In Vitro Delivery in COS, HT1080, and 293 Cell Lines

COS cells (available from the American Type Culture Collection, Rockville, MD, under Accession No. CRL 1651 and HT1080 cells (available from the American Type Culture Collection, Rockville, MD, under Accession No. CCL 121) were cultured and transfected with pCMVkmLUC and various polycationic agents of the present invention (described in Example 1) according to the transfection protocol described in Example 4. Luciferase activity was assayed according to the method described in Example 4. Total cell protein was measured using a Pierce BCA kit according to manufacturer's directions.

The results, shown in Figure 7A, indicate that the ability of the polycationic agents to mediate transfection is not dependent on cell line type. Polycationic agents having a repeating trimer motif of neutral and cationic sidechains were particularly effective at mediating transfection.

Transfection efficiencies for a homologous series of cationic peptoids were evaluated. Specifically, cationic peptoids RZ-110-1 (18-mer), RZ-120-3 (24mer), RZ120-7 (36mer), and RZ120-11 (48mer), which have the same repeating (HHP) motif were evaluated for their ability to transfect COS and HT1080 cells. These polycationic agents were complexed with pCMVkmLUC at a 2:1, + to - charge ratio. The concentration of negative charges on DNA was calculated using 3.03 nmol of phosphate per 1 µg of DNA, on the basis of the average molecular weight of 330 for each nucleotide. The formula weight of the polycationic agent was calculated as a semi-trifluoroacetate salt (50% of amino groups form salt with TFA), and the concentration of the polycationic agent was determined on the basis of the weight of the lyophilized peptoid. Amino groups were formally considered to be fully protonated to obtain the number of positive charges on the polycationic agent interest when calculating the + to - charge.

As shown in Figure 7B, transfection efficiencies for this particular series of cationic peptoids were largely independent of oligomer length for peptoids having 24 or more monomeric units.

Transfection efficiencies using polycationic agent RZ145-1 and commercially available cationic lipids, DMRIE-C[™], Lipofectin[®] and lipofectamine were evaluated. In these experiments RZ145-1 was complexed with pCMVkmLUC at a 2:1, + to - charge ratio. Transfection with DMRIE-C[™] Lipofectin[®], lipofectamine was conducted according to

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manufacturer's directions. The cationic lipids were also employed at a 2:1, + to - charge ratio. 293 human embryonic kidney cells (Microbix, Toronto, Ontario, Canada), HT1080 cells, and NIH-3T3 cells (available from the American Type Culture Collection, Rockville, MD, Accession No. CRL 1658) were transfected, cultured either in the presence or absence of 10% serum, then assayed for luciferase production using the same protocol as described in Example 4. Luciferase was measured, as described in Example 4, 48 hours after initial transfection. Total cell protein was measured using a Pierce BCA kit according to manufacturer's directions.

The results, shown in Figure 8, indicate that, in contrast to Lipofectin® and lipofectamine, which were respectively 10- and 100-fold less efficient in the presence of serum, gene transfer mediated by polycationic agent RZ145-1 was insensitive to the presence of serum.

Transfection mediated by polycationic polymers, such as polylysine and histones, is greatly enhanced by addition of chloroquine to the transfection media. To determine whether chloroquine affected transfection mediated by polycationic agents of the present invention, HT1080 and 293 cells were transfected using RZ145-1 in the presence and absence of chloroquine. As a control, the same cell lines were transfected with polylysine both in the presence and absence of chloroquine. The results, shown in Figure 9, indicate that the polycationic agent RZ145-1 was equally effective at mediating transfection both with and without chloroquine. In contrast, polylysine-mediated transfection in the absence of chloroquine was 100-fold lower than polylysine mediated transfection in the presence of chloroquine. In addition, the results indicate that cationic peptoid mediated transfection is more efficient than polylysine mediated transfection.

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Example 10

Preparation of a Stable Formulation of DNA/Polycationic Agent Complex

A. DNA/Polycatinic Agent Complex Formation (2:1, + to - Change Ratio)

All operations were carried out at ambient temperature. DGPW (diagnosis grade purified water) was used to prepare the stock solutions. Both the plycatinic agent and DNA samples had low salt concentrations (i.e., < 1mM) to avoid precipitation.

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(1) Batch Method

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Complexes of polycationic agent RZ145-1 and pCMVkmLUC, as follows, for up to 250 µg DNA. DNA (i.e., pCMVkmLUC) was diluted with 30% (v/v) ethanol in water to a concentration of 50 µg/ml corresponding to 151 µM of negative charge. RZ145-1 was diluted to 23.2 µM in 30% ethanol in water. To 1 part of the polycationic agent solution was added 1 part of DNA solution as quickly as possible with gentle agitation. The DNA solution was added to the solution of polycationic agent (rather than vice-versa) to avoid precipitation. Slow addition of the two solutions was avoided to avoid precipitation and the formation of large complexes.

(2) Continuous Method

For more than 250 µg of DNA, a continuous method for preparing concentrated formulations of polycationic agent/DNA complex is preferred. The DNA and peptoid solutions were prepared as above and placed into separate bottles. Each bottle was connected to one port of a mixing tee. The bottles were simultaneously pressurized with 2 to 3 psi to deliver the two streams to the mixing tee at the same flow rate (e.g., 20 ml/min or higher).

B. Concentration Step

Two milliliters of the DNA-polycationic agent complex from part A was placed in a Centricon*-100 (Amico Inc. Beverly, MA), and centrifuged at 1000 x g for 30 minutes or until the volume of the retentate containing polycationic agentDNA complex was approximately 50 µl. The filtrate was removed from the bottom receiver. The retentate was diluted with 2 ml of 5% glucose, and concentrated to 50 µl again. This operation was repeated again to produce a concentrated complex solution containing 1 mg/ml DNA in 5% glucose. This concentration step can be conducted at either 4°C or at ambient temperature. The ethanol content of the final concentrated solution was less than 0.1%. No precipitation was observed in the concentrated solution.

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Zuckermann et al.
- (ii) TITLE OF INVENTION: Compositions and Methods for Polynucleotide Delivery
- (iii) NUMBER OF SEQUENCES: 4
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Chiron Corporation
 - (B) STREET: 4560 Horton Street
 - (C) CITY: Emeryville
 - (D) STATE: California
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 94608-2916
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Fujita, Sharon M.
 - (B) REGISTRATION NUMBER: 38,459
 - (C) REFERENCE/DOCKET NUMBER: 1218.002
 - (ix) TELECOMMUNICATION INFORMATION:
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- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9600 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GCGGCCGCGG	AATTCTCATG	TTTGACAGCT	TATCATCGAT	AAGCTGATCC	TCACAGGCCG	60
CACCCAGCTT	TTCTTCCGTT	GCCCCAGTAG	CATCTCTGTC	TGGTGACCTT	GAAGAGGAAG	120
AGGAGGGGTC	CCGAGAATCC	CCATCCCTAC	CGTCCAGCAA	AAAGGGGGAC	GAGGAATTTG	180
AGGCCTGGCT	TGAGGCTCAG	GACGCAAATC	TTGAGGATGT	TCAGCGGGAG	TTTTCCGGGC	240
TGCGAGTAAT	TGGTGATGAG	GACGAGGATG	GTTCGGAGGA	TGGGGAATTT	TCAGACCTGG	300
ATCTGTCTGA	CAGCGACCAT	GAAGGGGATG	AGGGTGGGG	GGCTGTTGGA	GGGGGCAGGA	360
GTCTGCACTC	CCTGTATTCA	CTGAGCGTCG	TCTAATAAAG	ATGTCTATTG	ATCTCTTTTA	420
GTGTGAATCA	TGTCTGACGA	GGGGCCAGGT	ACAGGACCTG	GAAATGGCCT	AGGAGAGAAG	480
GGAGACACAT	CTGGACCAGA	AGGCTCCGGC	GGCAGTGGAC	CTCAAAGAAG	AGGGGGTGAT	540
AACCATGGAC	GAGGACGGG	AAGAGGACGA	GGACGAGGAG	GCGGAAGACC	AGGAGCCCCG	600
GGCGGCTCAG	GATCAGGGCC	AAGACATAGA	GATGGTGTCC	GGAGACCCCA	AAAACGTCCA	660
AGTTGCATTG	GCTGCAAAGG	GACCCACGGT	GGAACAGGAG	CAGGAGCAGG	AGCGGGAGGG	720

GCAGGAGCAG	GAGGGCAGG	AGCAGGAGGA	GGGGCAGGAG	CAGGAGGAGG	GGCAGGAGGG	780
GCAGGAGGGG	CAGGAGGGC	AGGAGCAGGA	GGAGGGCAG	GAGCAGGAGG	AGGGGCAGGA	840
				GAGGAGGGC		900
				CAGGAGGAGG		960
				GAGGGGCAGG		1020
				GAGGGCAGG		1080
				GAGCAGGAGG		1140
,				GAGGGGCAGG		1200
				CAGGAGGAGG		1260
				GGGCAGGAGG		1320
				CAGGAGCAGG		1380
				GAGGAGGCAG GAGGCCGCCG		1440
				GGAGAGGTCG		1500
				CCGGGTCTCC		1560 1620
				AAGCCGATTA		1680
				CGGGAGCGAT		1740
				GGGGTCAGGG		1800
				AAGGAGGTTC		1860
				GGAGTCACGT		1920
ACCGACGAAG	GAACTTGGGT	CGCCGGTGTG	TTCGTATATG	GAGGTAGTAA	GACCTCCCTT	1980
TACAACCTAA	GGCGAGGAAC	TGCCCTTGCT	ATTCCACAAT	GTCGTCTTAC	ACCATTGAGT	2040
CGTCTCCCCT	TTGGAATGGC	CCCTGGACCC	GGCCCACAAC	CTGGCCCGCT	AAGGGAGTCC	2100
ATTGTCTGTT	ATTTCATGGT	CTTTTTACAA	ACTCATATAT	TTGCTGAGGT	TTTGAAGGAT	2160
				GCAATATCAG		2220
				CACCTATGGT		2280
				GTGATGGAGA		2340
				ATCGTATTAA		2400
				CGTGCTGTTG		2460
				GGCAATTGGG		2520
				TTGGAAAACA GCCTCCTTAA		2580 2640
				AAAATTCACG		2700
				ACTGGGTATC		2760
				TTAGGATAGC		2820
				TTAGGATAGC		2880
				TTAGGATAGC		2940
CAGATATAGA	TTAGGATAGC	CTATGCTACC	CAGATATAGA	TTAGGATAGC	ATATGCTACC	3000
CAGATATAGA	TTAGGATAGC	ATATGCTATC	CAGATATTTG	GGTAGTATAT	GCTACCCAGA	3060
TATAAATTAG	GATAGCATAT	ACTACCCTAA	TCTCTATTAG	GATAGCATAT	GCTACCCGGA	3120
				GATAGCATAT		3180
				GATAGCATAT		3240
				GATAGCCTAT		3300
				GTATATGCTA		3360
				GACCAACAAC		3420
				CAGCAATCGC		3480 3540
				AAGTTATTAC		3600
				GTGCCCCCAC		3660
				ATTGGCGTGG		3720
				CTGTCGGCGT		3780
				GTCTTGGTCC		3840
CACATCTTAA	TAACCCCAGT	ATCATATTGC	ACTAGGATTA	TGTGTTGCCC	ATAGCCATAA	3900
ATTCGTGTGA	GATGGACATC	CAGTCTTTAC	GGCTTGTCCC	CACCCCATGG	ATTTCTATTG	3960
TTAAAGATAT	TCAGAATGTT	TCATTCCTAC	ACTAGTATTT	ATTGCCCAAG	GGGTTTGTGA	4020
				CCCCGTCCAA		4080
				ACACCTTACT		4140
				GCAGGCGAAG		4200
				GACTAAAATG		4260
				CTCATGGGGT		4320
				CATATGCTTC		4380
				CTACTACCCG		4440
GCIACCOGTT	AGGGTTAAC	ANGGGGGCCT		ATTGCTAATG	CCCTCTTGAG	4500
			78			

BNSDOCID: <WO__9806437A2_i_>

GGTCCGCTTA	TCGGTAGCTA	CACAGGCCCC	TCTGATTGAC	GTTGGTGTAG	CCTCCCGTAG	4560
TCTTCCTGGG	CCCCTGGGAG	GTACATGTCC	CCCAGCATTG	GTGTAAGAGC	TTCAGCCAAG	4620
AGTTACACAT	AAAGGCAATG	TTGTGTTGCA	GTCCACAGCA	TGCAAAGTCT	GCTCCAGGAT	4680
GAAAGCCACT	CAGTGTTGGC	AAATGTGCAC	ATCCATTTAT	AAGGATGTCA	ACTACAGTCA	4740
GAGAACCCCT	TTGTGTTTGG	TCCCCCCCG	TGTCACATGT	GGAACAGGGC	CCAGTTGGCA	4800
AGTTGTACCA	ACCAACTGAA	GGGATTACAT	GCACTGCCCC	GAATACAAAA	CAAAAGCGCT	4860
CCTCGTACCA	GCGAAGAAGG	GGCAGAGATG	CCGTAGTCAG	GTTTAGTTCG	TCCGGCGGCG	4920
GGGCGATCCG	CGGCCGCCTG	CGCGCTCGCT	CGCTCACTGA	GGCCGCCCGG	GCAAAGCCCG	4980
			CAGTGAGCGA			5040
			AGTTAATGAT			5100
			TTGCATACGT			5160
			CCGCCATGTT			5220
			GTTCATAGCC			
			TGACCGCCCA			5280
						5340
			CCAATAGGGA			5400
			GCAGTACATC			5460
			TGGCCCGCCT			5520
			ATCTACGTAT			5580
			CGTGGATAGC			5640
			AGTTTGTTTT			5700
			TTGACGCAAA			5760
			GTGAACCGTC			5820
			CGGGACCGAT			5880
			AAGAGTGACG			5940
			GCTATACTGT			6000
			ATAGCTTAGC			6060
CCATTATTGA	CCACTCCCCT	ATTGGTGACG	ATACTTTCCA	TTACTAATCC	ATAACATGGC	6120
TCTTTGCCAC	AACTATCTCT	ATTGGCTATA	TGCCAATACT	CTGTCCTTCA	GAGACTGACA	6180
CGGACTCTGT	ATTTTTACAG	GATGGGGTCC	ATTTATTATT	TACAAATTCA	CATATACAAC	6240
AACGCCGTCC	CCCGTGCCCG	CAGTTTTTAT	TAAACATAGC	GTGGGATCTC	CGACATCTCG	6300
GGTACGTGTT	CCGGACATGG	GCTCTTCTCC	GGTAGCGGCG	GAGCTTCCAC	ATCCGAGCCC	6360
TGGTCCCATC	CGTCCAGCGG	CTCATGGTCG	CTCGGCAGCT	CCTTGCTCCT	AACAGTGGAG	6420
GCCAGACTTA	GGCACAGCAC	AATGCCCACC	ACCACCAGTG	TGCCGCACAA	GGCCGTGGCG	6480
GTAGGGTATG	TGTCTGAAAA	TGAGCTCGGA	GATTGGGCTC	GCACCTGGAC	GCAGATGGAA	6540
GACTTAAGGC	AGCGGCAGAA	GAAGATGCAG	GCAGCTGAGT	TGTTGTATTC	TGATAAGAGT	6600
CAGAGGTAAC	TCCCGTTGCG	GTGCTGTTAA	CGGTGGAGGG	CAGTGTAGTC	TGAGCAGTAC	6660
TCGTTGCTGC	CGCGCGCGCC	ACCAGACATA	ATAGCTGACA	GACTAACAGA	CTGTTCCTTT	6720
CCATGGGTCT	TTTCTGCAGT	CACCGTCGTC	GACCTAAGAA	TTCAGACTCG	AGCAAGTCTA	6780
GAAAGCCATG	GATATCGGAT	CCACTACGCG	TTAGAGCTCG	CTGATCAGCC	TCGACTGTGC	6840
CTTCTAGTTG	CCAGCCATCT	GTTGTTTGCC	CCTCCCCCGT	GCCTTCCTTG	ACCCTGGAAG	6900
GTGCCACTCC	CACTGTCCTT	TCCTAATAAA	ATGAGGAAAT	TGCATCGCAT	TGTCTGAGTA	6960
GGTGTCATTC	TATTCTGGGG	GGTGGGGTGG	GGCAGGACAG	CAAGGGGGAG	GATTGGGAAG	7020
ACAATAGCAG	GGGGGTGGGC	GAAGAACTCC	AGCATGAGAT	CCCCGCGCTG	GAGGATCATC	7080
CAGCCGGCGT	CCCGGAAAAC	GATTCCGAAG	CCCAACCTTT	CATAGAAGGC	GGCGGTGGAA	7140
TCGAAATCTC	GTGATGGCAG	GTTGGGCGTC	GCTTGGTCGG	TCATTTCGGT	AGATAAGTAG	7200
CATGGCGGGT	TAATCATTAA	CTACAAGGAA	CCCCTAGTGA	TGGAGTTGGC	CACTCCCTCT	7260
CTGCGCGCTC	GCTCGCTCAC	TGAGGCCGGG	CGACCAAAGG	TCGCCCGACG	CCCGGGCTTT	7320
GCCCGGGCGG	CCTCAGTGAG	CGAGCGAGCG	CGCAGCGAAC	CCCAGAGTCC	CGCTCAGAAG	7380
AACTCGTCAA	GAAGGCGATA	GAAGGCGATG	CGCTGCGAAT	CGGGAGCGGC	GATACCGTAA	7440
AGCACGAGGA	AGCGGTCAGC	CCATTCGCCG	CCAAGCTCTT	CAGCAATATC	ACGGGTAGCC	7500
AACGCTATGT	CCTGATAGCG	GTCCGCCACA	CCCAGCCGGC	CACAGTCGAT	GAATCCAGAA	7560
AAGCGGCCAT	TTTCCACCAT	GATATTCGGC	AAGCAGGCAT	CGCCATGGGT	CACGACGAGA	7620
TCCTCGCCGT	CGGGCATGCG	CGCCTTGAGC	CTGGCGAACA	GTTCGGCTGG	CGCGAGCCCC	7680
TGATGCTCTT	CGTCCAGATC	ATCCTGATCG	ACAAGACCGG	CTTCCATCCG	AGTACGTGCT	7740
CGCTCGATGC	GATGTTTCGC	TTGGTGGTCG	AATGGGCAGG	TAGCCGGATC	AAGCGTATGC	7800
AGCCGCCGCA	TTGCATCAGC	CATGATGGAT	ACTTTCTCGG	CAGGAGCAAG	GTGAGATGAC	7860
AGGAGATCCT	GCCCCGGCAC	TTCGCCCAAT	AGCAGCCAGT	CCCTTCCCGC	TTCAGTGACA	7920
			GTCGTGGCCA		· -	7980
			AGGTCGGTCT			8040
			TCAGAGCAGC			8100
			GCCGGAGAAC			8160
			TCTTGATCAG			8220
			TTTACTTTGC			8280
			79			

GAGGGCGCCC CAGCTGGCAA TTCCGGTTCG CTTGCTGTCC ATAAAACCGC CCAGTCTAGC 8340 TATCGCCATG TAAGCCCACT GCAAGCTACC TGCTTTCTCT TTGCGCTTGC GTTTTCCCTT GTCCAGATAG CCCAGTAGCT GACATTCATC CGGGGTCAGC ACCGTTTCTG CGGACTGGCT 8460 TTCTACGTGT TCCGCTTCCT TTAGCAGCCC TTGCGCCCTG AGTGCTTGCG GCAGCGTGAA GCTGTCAATT CCGCGTTAAA TTTTTGTTAA ATCAGCTCAT TTTTTAACCA ATAGGCCGAA ATCGGCAAAA TCCCTTATAA ATCAAAAGAA TAGCCCGAGA TAGGGTTGAG TGTTGTTCCA GTTTGGAACA AGAGTCCACT ATTAAAGAAC GTGGACTCCA ACGTCAAAGG GCGAAAAACC GTCTATCAGG GCGATGGCGG ATCAGCTTAT GCGGTGTGAA ATACCGCACA GATGCGTAAG 8760 GAGAAAATAC CGCATCAGGC GCTCTTCCGC TTCCTCGCTC ACTGACTCGC TGCGCTCGGT 8820 CGTTCGGCTG CGGCGAGCGG TATCAGCTCA CTCAAAGGCG GTAATACGGT TATCCACAGA ATCAGGGGAT AACGCAGGAA AGAACATGTG AGCAAAAGGC CAGCAAAAGG CCAGGAACCG TAAAAAGGCC GCGTTGCTGG CGTTTTTCCA TAGGCTCCGC CCCCCTGACG AGCATCACAA AAATCGACGC TCAAGTCAGA GGTGGCGAAA CCCGACAGGA CTATAAAGAT ACCAGGCGTT TCCCCCTGGA AGCTCCCTCG TGCGCTCTCC TGTTCCGACC CTGCCGCTTA CCGGATACCT 9120 GTCCGCCTTT CTCCCTTCGG GAAGCGTGGC GCTTTCTCAT AGCTCACGCT GTAGGTATCT 9180 CAGTTCGGTG TAGGTCGTTC GCTCCAAGCT GGGCTGTGTG CACGAACCCC CCGTTCAGCC 9240 CGACCGCTGC GCCTTATCCG GTAACTATCG TCTTGAGTCC AACCCGGTAA GACACGACTT 9300 ATCGCCACTG GCAGCAGCCA CTGGTAACAG GATTAGCAGA GCGAGGTATG TAGGCGGTGC 9360 TACAGAGTTC TTGAAGTGGT GGCCTAACTA CGGCTACACT AGAAGGACAG TATTTGGTAT 9420 CTGCGCTCTG CTGAAGCCAG TTACCTTCGG AAAAAGAGTT GGTAGCTCTT GATCCGGCAA 9480 ACAAACCACC GCTGGTAGCG GCGGTTTTTT GTTTGCAAGC AGCAGATTAC GCGCAGAAAA 9540 AAAGGATCTC AAGAAGATCC TTTGATCTTT TCTTACTGAA CGGTGATCCC CACCGGAATT 9600

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4328 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

GCCGCGGAAT TTCGACTCTA GGCCATTGCA TACGTTGTAT CTATATCATA ATATGTACAT 60 TTATATTGGC TCATGTCCAA TATGACCGCC ATGTTGACAT TGATTATTGA CTAGTTATTA 120 ATAGTAATCA ATTACGGGGT CATTAGTTCA TAGCCCATAT ATGGAGTTCC GCGTTACATA 180 ACTTACGGTA AATGGCCCGC CTGGCTGACC GCCCAACGAC CCCCGCCCAT TGACGTCAAT AATGACGTAT GTTCCCATAG TAACGCCAAT AGGGACTTTC CATTGACGTC AATGGGTGGA GTATTTACGG TAAACTGCCC ACTTGGCAGT ACATCAAGTG TATCATATGC CAAGTCCGCC 360 CCCTATTGAC GTCAATGACG GTAAATGGCC CGCCTGGCAT TATGCCCAGT ACATGACCTT 420 ACGGGACTTT CCTACTTGGC AGTACATCTA CGTATTAGTC ATCGCTATTA CCATGGTGAT 480 GCGGTTTTGG CAGTACACCA ATGGGCGTGG ATAGCGGTTT GACTCACGGG GATTTCCAAG TCTCCACCCC ATTGACGTCA ATGGGAGTTT GTTTTGGCAC CAAAATCAAC GGGACTTTCC 600 AAAATGTCGT AATAACCCCG CCCCGTTGAC GCAAATGGGC GGTAGGCGTG TACGGTGGGA 660 GGTCTATATA AGCAGAGCTC GTTTAGTGAA CCGTCAGATC GCCTGGAGAC GCCATCCACG 720 CTGTTTTGAC CTCCATAGAA GACACCGGGA CCGATCCAGC CTCCGCGGCC GGGAACGGTG 780 CATTGGAACG CGGATTCCCC GTGCCAAGAG TGACGTAAGT ACCGCCTATA GACTCTATAG GCACACCCCT TTGGCTCTTA TGCATGCTAT ACTGTTTTTG GCTTGGGGCC TATACACCCC 900 CGCTCCTTAT GCTATAGGTG ATGGTATAGC TTAGCCTATA GGTGTGGGTT ATTGACCATT 960 ATTGACCACT CCCCTATTGG TGACGATACT TTCCATTACT AATCCATAAC ATGGCTCTTT 1020 GCCACAACTA TCTCTATTGG CTATATGCCA ATACTCTGTC CTTCAGAGAC TGACACGGAC TCTGTATTTT TACAGGATGG GGTCCATTTA TTATTTACAA ATTCACATAT ACAACAACGC 1140 CGTCCCCCGT GCCCGCAGTT TTTATTAAAC ATAGCGTGGG ATCTCCGACA TCTCGGGTAC 1200 GTGTTCCGGA CATGGGCTCT TCTCCGGTAG CGGCGGAGCT TCCACATCCG AGCCCTGGTC 1260 CCATCCGTCC AGCGGCTCAT GGTCGCTCGG CAGCTCCTTG CTCCTAACAG TGGAGGCCAG 1320 ACTTAGGCAC AGCACAATGC CCACCACCAC CAGTGTGCCG CACAAGGCCG TGGCGGTAGG 1380 GTATGTGTCT GAAAATGAGC TCGGAGATTG GGCTCGCACC TGGACGCAGA TGGAAGACTT AAGGCAGCGG CAGAAGAAGA TGCAGGCAGC TGAGTTGTTG TATTCTGATA AGAGTCAGAG GTAACTCCCG TTGCGGTGCT GTTAACGGTG GAGGGCAGTG TAGTCTGAGC AGTACTCGTT 1560 GCTGCCGCGC GCGCCACCAG ACATAATAGC TGACAGACTA ACAGACTGTT CCTTTCCATG 1620 GGTCTTTTCT GCAGTCACCG TCGTCGACCT AAGAATTCAG ACTCGAGCAA GTCTAGAAAG 1680 CCATGGATAT CGGATCCACT ACGCGTTAGA GCTCGCTGAT CAGCCTCGAC TGTGCCTTCT 1740 AGTTGCCAGC CATCTGTTGT TTGCCCCTCC CCCGTGCCTT CCTTGACCCT GGAAGGTGCC 1800 ACTCCCACTG TCCTTTCCTA ATAAAATGAG GAAATTGCAT CGCATTGTCT GAGTAGGTGT 1860

		GGTGGGGCAG				1920
		ACTCCAGCAT				1980
GGCGTCCCGG	AAAACGATTC	CGAAGCCCAA	CCTTTCATAG	AAGGCGGCGG	TGGAATCGAA	2040
ATCTCGTGAT	GGCAGGTTGG	GCGTCGCTTG	GTCGGTCATT	TCGAACCCCA	GAGTCCCGCT	2100
CAGAAGAACT	CGTCAAGAAG	GCGATAGAAG	GCGATGCGCT	GCGAATCGGG	AGCGGCGATA	2160
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GTAGCCAACG	CTATGTCCTG	ATAGCGGTCC	GCCACACCCA	GCCGGCCACA	GTCGATGAAT	2280
CCAGAAAAGC	GGCCATTTTC	CACCATGATA	TTCGGCAAGC	AGGCATCGCC	ATGGGTCACG	2340
ACGAGATCCT	CGCCGTCGGG	CATGCGCGCC	TTGAGCCTGG	CGAACAGTTC	GGCTGGCGCG	2400
AGCCCCTGAT	GCTCTTCGTC	CAGATCATCC	TGATCGACAA	GACCGGCTTC	CATCCGAGTA	2460
CGTGCTCGCT	CGATGCGATG	TTTCGCTTGG	TGGTCGAATG	GGCAGGTAGC	CGGATCAAGC	2520
GTATGCAGCC	GCCGCATTGC	ATCAGCCATG	ATGGATACTT	TCTCGGCAGG	AGCAAGGTGA	2580
GATGACAGGA	GATCCTGCCC	CGGCACTTCG	CCCAATAGCA	GCCAGTCCCT	TCCCGCTTCA	2640
GTGACAACGT	CGAGCACAGC	TGCGCAAGGA	ACGCCCGTCG	TGGCCAGCCA	CGATAGCCGC	2700
GCTGCCTCGT	CCTGCAGTTC	ATTCAGGGCA	CCGGACAGGT	CGGTCTTGAC	AAAAAGAACC	2760
GGCGCCCCT	GCGCTGACAG	CCGGAACACG	GCGGCATCAG	AGCAGCCGAT	TGTCTGTTGT	2820
GCCCAGTCAT	AGCCGAATAG	CCTCTCCACC	CAAGCGGCCG	GAGAACCTGC	GTGCAATCCA	2880
TCTTGTTCAA	TCATGCGAAA	CGATCCTCAT	CCTGTCTCTT	GATCAGATCT	TGATCCCCTG	2940
CGCCATCAGA	TCCTTGGCGG	CAAGAAAGCC	ATCCAGTTTA	CTTTGCAGGG	CTTCCCAACC	3000
TTACCAGAGG	GCGCCCCAGC	TGGCAATTCC	GGTTCGCTTG	CTGTCCATAA	AACCGCCCAG	3060
TCTAGCTATC	GCCATGTAAG	CCCACTGCAA	GCTACCTGCT	TTCTCTTTGC	GCTTGCGTTT	3120
TCCCTTGTCC	AGATAGCCCA	GTAGCTGACA	TTCATCCGGG	GTCAGCACCG	TTTCTGCGGA	3180
CTGGCTTTCT	ACGTGTTCCG	CTTCCTTTAG	CAGCCCTTGC	GCCCTGAGTG	CTTGCGGCAG	3240
CGTGAAGCTG	TCAATTCCGC	GTTAAATTTT	TGTTAAATCA	GCTCATTTTT	TAACCAATAG	3300
GCCGAAATCG	GCAAAATCCC	TTATAAATCA	AAAGAATAGC	CCGAGATAGG	GTTGAGTGTT	3360
GTTCCAGTTT	GGAACAAGAG	TCCACTATTA	AAGAACGTGG	ACTCCAACGT	CAAAGGGCGA	3420
AAAACCGTCT	ATCAGGGCGA	TGGCGGATCA	GCTTATGCGG	TGTGAAATAC	CGCACAGATG	3480
CGTAAGGAGA	AAATACCGCA	TCAGGCGCTC	TTCCGCTTCC	TCGCTCACTG	ACTCGCTGCG	3540
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CACAGAATCA	GGGGATAACG	CAGGAAAGAA	CATGTGAGCA	AAAGGCCAGC	AAAAGGCCAG	3660
GAACCGTAAA	AAGGCCGCGT	TGCTGGCGTT	TTTCCATAGG	CTCCGCCCCC	CTGACGAGCA	3720
TCACAAAAAT	CGACGCTCAA	GTCAGAGGTG	GCGAAACCCG	ACAGGACTAT	AAAGATACCA	3780
GGCGTTTCCC	CCTGGAAGCT	CCCTCGTGCG	CTCTCCTGTT	CCGACCCTGC	CGCTTACCGG	3840
ATACCTGTCC	GCCTTTCTCC	CTTCGGGAAG	CGTGGCGCTT	TCTCATAGCT	CACGCTGTAG	3900
GTATCTCAGT	TCGGTGTAGG	TCGTTCGCTC	CAAGCTGGGC	TGTGTGCACG	AACCCCCCGT	3960
TCAGCCCGAC	CGCTGCGCCT	TATCCGGTAA	CTATCGTCTT	GAGTCCAACC	CGGTAAGACA	4020
		CAGCCACTGG				4080
		AGTGGTGGCC				4140
		AGCCAGTTAC				4200
		GTAGCGGCGG				4260
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GAATTGCG						4328

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5107 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

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TTAATAGTAA	TCAATTACGG	GGTCATTAGT	TCATAGCCCA	TATATGGAGT	TCCGCGTTAC	180
ATAACTTACG	GTAAATGGCC	CGCCTGGCTG	ACCGCCCAAC	GACCCCCGCC	CATTGACGTC	240
AATAATGACG	TATGTTCCCA	TAGTAACGCC	AATAGGGACT	TTCCATTGAC	GTCAATGGGT	300
GGAGTATTTA	CGGTAAACTG	CCCACTTGGC	AGTACATCAA	GTGTATCATA	TGCCAAGTCC	360
GCCCCTATT	GACGTCAATG	ACGGTAAATG	GCCCGCCTGG	CATTATGCCC	AGTACATGAC	420
CTTACGGGAC	TTTCCTACTT	GGCAGTACAT	CTACGTATTA	GTCATCGCTA	TTACCATGGT	480
GATGCGGTTT	TGGCAGTACA	CCAATGGGCG	TGGATAGCGG	TTTGACTCAC	GGGGATTTCC	540
AAGTCTCCAC	CCCATTGACG	TCAATGGGAG	TTTGTTTTGG	CACCAAAATC	AACGGGACTT	600
TCCAAAATGT	CGTAATAACC	CCGCCCCGTT	GACGCAAATG	GGCGGTAGGC	GTGTACGGTG	660

GGAGGTCTAT	ATAAGCAGAG	CTCGTTTAGT	GAACCGTCAG	ATCGCCTGGA	GACGCCATCC	720
ACGCTGTTTT	GACCTCCATA	GAAGACACCG	GGACCGATCC	AGCCTCCGCG	GCCGGGAACG	780
	ACGCGGATTC					840
TAGGCACACC	CCTTTGGCTC	TTATGCATGC	TATACTGTTT	TTGGCTTGGG	GCCTATACAC	900
CCCCGCTCCT	TATGCTATAG	GTGATGGTAT	AGCTTAGCCT	ATAGGTGTGG	GTTATTGACC	960
ATTATTGACC	ACTCCCCTAT	TGGTGACGAT	ACTTTCCATT	ACTAATCCAT	AACATGGCTC	1020
	CTATCTCTAT					1080
	TTTTACAGGA					1140
	CGTGCCCGCA					1200
	GGACATGGGC					1260
	TCCAGCGGCT					1320
	CACAGCACAA					1320
	TCTGAAAATG					
	CGGCAGAAGA					1440
	CCGTTGCGGT					1500
	CGCGCGCCAC					1560
	TCTGCAGTCA					1620
	ACCCCGGCCA					1680
	GTCTCTCGTG					1740
						1800
	TGACAGCCGA					1860
	GGGCTGTTCC					1920
	CTTCTATGCC					1980
	GGCCCTGCTC					2040
	TTTCGAGCCC					2100
	TCTGCTTCGG					2160
	TCCACTCCGA					2220
	CCTCCGGGGA					2280
	AGGTGCGTCC					2340
	TCCCTCACCA					2400
	GACACTCCAG					2460
	TGAGATCCAC					2520
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	GTCCTTTCCT					2640
	CTGGGGGGTG					2700
	GTGGGCGAAG					2760
	GAAAACGATT					2820
	TGGCAGGTTG					2880
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	ACGAGGAAGC					3000
	GCTATGTCCT					3060
	CGGCCATTTT					3120
	TCGCCGTCGG					3180
	TGCTCTTCGT					3240
	TCGATGCGAT					3300
	CGCCGCATTG					3360
	AGATCCTGCC					3420
	TCGAGCACAG					3480
	TCCTGCAGTT					3540
	TGCGCTGACA					3600
	TAGCCGAATA					3660
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	ATCCTTGGCG					3780
	GGCGCCCCAG					3840
	CGCCATGTAA					3900
	CAGATAGCCC					3960
	TACGTGTTCC					4020
	GTCAATTCCG					4080
	GGCAAAATCC					4140
	TGGAACAAGA					4200
	TATCAGGGCG					4260
GCGTAAGGAG	AAAATACCGC	ATCAGGCGCT	CTTCCGCTTC	CTCGCTCACT	GACTCGCTGC	4320
GCTCGGTCGT	TCGGCTGCGG	CGAGCGGTAT	CAGCTCACTC	AAAGGCGGTA	ATACGGTTAT	4380
	AGGGGATAAC		ACATGTGAGC			4440
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BNSDOCID: <WO___9806437A2_I_>

GGAACCGTAA AAAGGCCGCG	TTGCTGGCGT	TTTTCCATAG	GCTCCGCCCC	CCTGACGAGC	4500
ATCACAAAAA TCGACGCTCA	AGTCAGAGGT	GGCGAAACCC	GACAGGACTA	TAAAGATACC	4560
AGGCGTTTCC CCCTGGAAGC	TCCCTCGTGC	GCTCTCCTGT	TCCGACCCTG	CCGCTTACCG	4620
GATACCTGTC CGCCTTTCTC	CCTTCGGGAA	GCGTGGCGCT	TTCTCATAGC	TCACGCTGTA	4680
GGTATCTCAG TTCGGTGTAG	GTCGTTCGCT	CCAAGCTGGG	CTGTGTGCAC	GAACCCCCCG	4740
TTCAGCCCGA CCGCTGCGCC	TTATCCGGTA	ACTATCGTCT	TGAGTCCAAC	CCGGTAAGAC	4800
ACGACTTATC GCCACTGGCA	GCAGCCACTG	GTAACAGGAT	TAGCAGAGCG	AGGTATGTAG	4860
GCGGTGCTAC AGAGTTCTTG	AAGTGGTGGC	CTAACTACGG	CTACACTAGA	AGGACAGTAT	4920
TTGGTATCTG CGCTCTGCTG	AAGCCAGTTA	CCTTCGGAAA	AAGAGTTGGT	AGCTCTTGAT	4980
CCGGCAAACA AACCACCGCT	GGTAGCGGCG	GTTTTTTGTT	TGCAAGCAGC	AGATTACGCG	5040
CAGAAAAAA GGATCTCAAG	AAGATCCTTT	GATCTTTTCT	TACTGAACGG	TGATCCCCAC	5100
CGGAATT					5107

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4818 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

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GCGGCCGCGG	AATTTCGACT	CTAGGCCATT	GCATACGTTG	TATCTATATC	ATAATATGTA	60
CATTTATATT	GGCTCATGTC	CAATATGACC	GCCATGTTGA	CATTGATTAT	TGACTAGTTA	120
TTAATAGTAA	TCAATTACGG	GGTCATTAGT	TCATAGCCCA	TATATGGAGT	TCCGCGTTAC	180
ATAACTTACG	GTAAATGGCC	CGCCTGGCTG	ACCGCCCAAC	GACCCCCGCC	CATTGACGTC	240
AATAATGACG	TATGTTCCCA	TAGTAACGCC	AATAGGGACT	TTCCATTGAC	GTCAATGGGT	300
GGAGTATTTA	CGGTAAACTG	CCCACTTGGC	AGTACATCAA	GTGTATCATA	TGCCAAGTCC	360
GCCCCTATT	GACGTCAATG	ACGGTAAATG	GCCCGCCTGG	CATTATGCCC	AGTACATGAC	420
CTTACGGGAC	TTTCCTACTT	GGCAGTACAT	CTACGTATTA	GTCATCGCTA	TTACCATGGT	480
GATGCGGTTT	TGGCAGTACA	CCAATGGGCG	TGGATAGCGG	TTTGACTCAC	GGGGATTTCC	540
AAGTCTCCAC	CCCATTGACG	TCAATGGGAG	TTTGTTTTGG	CACCAAAATC	AACGGGACTT	600
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ACGCTGTTTT	GACCTCCATA	GAAGACACCG	GGACCGATCC	AGCCTCCGCG	GCCGGGAACG	780
GTGCATTGGA	ACGCGGATTC	CCCGTGCCAA	GAGTGACGTA	AGTACCGCCT	ATAGACTCTA	840
TAGGCACACC	CCTTTGGCTC	TTATGCATGC	TATACTGTTT	TTGGCTTGGG	GCCTATACAC	900
CCCCGCTCCT	TATGCTATAG	GTGATGGTAT	AGCTTAGCCT	ATAGGTGTGG	GTTATTGACC	960
ATTATTGACC	ACTCCCCTAT	TGGTGACGAT	ACTTTCCATT	ACTAATCCAT	AACATGGCTC	1020
TTTGCCACAA	CTATCTCTAT	TGGCTATATG	CCAATACTCT	GTCCTTCAGA	GACTGACACG	1080 .
GACTCTGTAT	TTTTACAGGA	TGGGGTCCAT	TTATTATTTA	CAAATTCACA	TATACAACAA	1140
CGCCGTCCCC	CGTGCCCGCA	GTTTTTATTA	AACATAGCGT	GGGATCTCCG	ACATCTCGGG	1200
TACGTGTTCC	GGACATGGGC	TCTTCTCCGG	TAGCGGCGGA	GCTTCCACAT	CCGAGCCCTG	1260
GTCCCATCCG	TCCAGCGGCT	CATGGTCGCT	CGGCAGCTCC	TTGCTCCTAA	CAGTGGAGGC	1320
CAGACTTAGG	CACAGCACAA	TGCCCACCAC	CACCAGTGTG	CCGCACAAGG	CCGTGGCGGT	1380
AGGGTATGTG	TCTGAAAATG	AGCTCGGAGA	TTGGGCTCGC	ACCTGGACGC	AGATGGAAGA	1440
CTTAAGGCAG	CGGCAGAAGA	AGATGCAGGC	AGCTGAGTTG	TTGTATTCTG	ATAAGAGTCA	1500
GAGGTAACTC	CCGTTGCGGT	GCTGTTAACG	GTGGAGGGCA	GTGTAGTCTG	AGCAGTACTC	1560
GTTGCTGCCG	CGCGCGCCAC	CAGACATAAT	AGCTGACAGA	CTAACAGACT	GTTCCTTTCC	1620
ATGGGTCTTT	TCTGCAGTCA	CCGTCGTCGA	CCTAAGAATT	CAGACTCGAG	CAAGTCTAGA	1680
ATGTGCTGGA	GACCCCTGTG	TCGGTTCCTG	TGGCTTTGGT	CCTATCTGTC	TTATGTTCAA	1740
GCAGTGCCTA	TCCAGAAAGT	CCAGGATGAC	ACCAAAACCC	TCATCAAGAC	CATTGTCACC	1800
AGGATCAATG	ACATTTCACA	CACGCAGTCG	GTATCCGCCA	AGCAGAGGGT	CACTGGCTTG	1860
GACTTCATTC	CTGGGCTTCA	CCCCATTCTG	AGTTTGTCCA	AGATGGACCA	GACTCTGGCA	1920
GTCTATCAAC	AGGTCCTCAC	CAGCCTGCCT	TCCCAAAATG	TGCTGCAGAT	AGCCAATGAC	1980
CTGGAGAATC	TCCGAGACCT	CCTCCATCTG	CTGGCCTTCT	CCAAGAGCTG	CTCCCTGCCT	2040
CAGACCAGTG	GCCTGCAGAA	GCCAGAGAGC	CTGGATGGCG	TCCTGGAAGC	CTCACTCTAC	2100
TCCACAGAGG	TGGTGGCTTT	GAGCAGGCTG	CAGGGCTCTC	TGCAGGACAT	TCTTCAACAG	2160
TTGGATGTTA	GCCCTGAATG	TGAGGATCCA	CTACGCGTTA	GAGCTCGCTG	ATCAGCCTCG	2220
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${\tt CTGGAAGGTG}$	CCACTCCCAC	TGTCCTTTCC	TAATAAAATG	AGGAAATTGC	ATCGCATTGT	2340
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TCGGCTGGCG	CGAGCCCCTG	ATGCTCTTCG	TCCAGATCAT	CCTGATCGAC	AAGACCGGCT	2940
TCCATCCGAG	TACGTGCTCG	CTCGATGCGA	TGTTTCGCTT	GGTGGTCGAA	TGGGCAGGTA	3000
GCCGGATCAA	GCGTATGCAG	CCGCCGCATT	GCATCAGCCA	TGATGGATAC	TTTCTCGGCA	3060
GGAGCAAGGT	GAGATGACAG	GAGATCCTGC	CCCGGCACTT	CGCCCAATAG	CAGCCAGTCC	3120
CTTCCCGCTT	CAGTGACAAC	GTCGAGCACA	GCTGCGCAAG	GAACGCCCGT	CGTGGCCAGC	3180
CACGATAGCC	GCGCTGCCTC	GTCCTGCAGT	TCATTCAGGG	CACCGGACAG	GTCGGTCTTG	3240
ACAAAAAGAA	CCGGGCGCCC	CTGCGCTGAC	AGCCGGAACA	CGGCGGCATC	AGAGCAGCCG	3300
ATTGTCTGTT	GTGCCCAGTC	ATAGCCGAAT	AGCCTCTCCA	CCCAAGCGGC	CGGAGAACCT	3360
GCGTGCAATC	CATCTTGTTC	AATCATGCGA	AACGATCCTC	ATCCTGTCTC	TTGATCAGAT	3420
CTTGATCCCC	TGCGCCATCA	GATCCTTGGC	GGCAAGAAAG	CCATCCAGTT	TACTTTGCAG	3480
GGCTTCCCAA	CCTTACCAGA	GGGCGCCCA	GCTGGCAATT	CCGGTTCGCT	TGCTGTCCAT	3540
AAAACCGCCC	AGTCTAGCTA	TCGCCATGTA	AGCCCACTGC	AAGCTACCTG	CTTTCTCTTT	3600
GCGCTTGCGT	TTTCCCTTGT	CCAGATAGCC	CAGTAGCTGA	CATTCATCCG	GGGTCAGCAC	3660
CGTTTCTGCG	GACTGGCTTT	CTACGTGTTC	CGCTTCCTTT	AGCAGCCCTT	GCGCCCTGAG	3720
TGCTTGCGGC	AGCGTGAAGC	TGTCAATTCC	GCGTTAAATT	TTTGTTAAAT	CAGCTCATTT	3780
TTTAACCAAT	AGGCCGAAAT	CGGCAAAATC	CCTTATAAAT	CAAAAGAATA	GCCCGAGATA	3840
GGGTTGAGTG	TTGTTCCAGT	TTGGAACAAG	AGTCCACTAT	TAAAGAACGT	GGACTCCAAC	3900
GTCAAAGGGC	GAAAAACCGT	CTATCAGGGC	GATGGCGGAT	CAGCTTATGC	GGTGTGAAAT	3960
ACCGCACAGA	TGCGTAAGGA	GAAAATACCG	CATCAGGCGC	TCTTCCGCTT	CCTCGCTCAC	4020
TGACTCGCTG	CGCTCGGTCG	TTCGGCTGCG	GCGAGCGGTA	TCAGCTCACT	CAAAGGCGGT	4080
AATACGGTTA	TCCACAGAAT	CAGGGGATAA	CGCAGGAAAG	AACATGTGAG	CAAAAGGCCA	4140
GCAAAAGGCC	AGGAACCGTA	AAAAGGCCGC	GTTGCTGGCG	TTTTTCCATA	GGCTCCGCCC	4200
CCCTGACGAG	CATCACAAAA	ATCGACGCTC	AAGTCAGAGG	TGGCGAAACC	CGACAGGACT	4260
ATAAAGATAC	CAGGCGTTTC	CCCCTGGAAG	CTCCCTCGTG	CGCTCTCCTG	TTCCGACCCT	4320
			CCCTTCGGGA			4380
			GGTCGTTCGC			4440
			CTTATCCGGT			4500
			AGCAGCCACT			4560
		· ·	GAAGTGGTGG			4620
			GAAGCCAGTT			4680
			TGGTAGCGGC			4740
		AGGATCTCAA	GAAGATCCTT	TGATCTTTTC	TTACTGAACG	4800
GTGATCCCCA	CCGGAATT					4818

WHAT IS CLAIMED:

1. A polycationic agent having the following formula:

$$Ta - \begin{bmatrix} R_1 & R_2 & O \\ -N - C - C - C - C \\ \frac{1}{R_3} & R_3 \end{bmatrix} - Tc$$

wherein

n is an integer from 10 to 100;

R, R, and R, for each monomer,

are independently selected from moieties having a molecular weight from 1 to 200 daltons;

Ta and Tc are terminating groups;

R, is not hydrogen for at least one monomer;

wherein said polycationic agent comprises at least 25% positively charged monomers, excluding the terminal groups, and

wherein said polycationic agent exhibits a net positive electrical charge at a physiological pH.

- 2. The polycationic agent according to claim 1, wherein said polycationic agent comprises repeating trimers.
- 3. The polycationic agent according to claim 2, wherein two R₁ groups in each trimer are neutral moieties and one R₁ group in each trimer is a cationic moiety.
- 4. The polycationic agent according to claim 1, wherein R₁ is selected from the group consisting of positively charged, negatively charged and neutral moieties.
- 5. The polycationic agent according to claim 1, wherein R_1 is selected from substituents found on amino acids.

6. The polycationic agent of claim 1, wherein R₁ is selected from the group consisting of aromatic and aliphatic groups.

- 7. The polycationic agent according to claim 1, wherein at least one R₁ is selected from the group consisting of alkylammonium, aminoalkyl, guanidinoalkyl, amidinoalkyl, aminocyclohexyl, piperidyl, guanidinobenzyl, amidinobenzyl, pyridylmethyl, aminobenzyl, alkyphenyl, indolylalkyl, alkoxyphenylalkyl, halophenylalkyl, and hydroxyphenylalkyl.
- 8. The polycationic agent according to claim 3, wherein said cationic moeity is aminoethyl.
- 9. The polycationic agent according to claim 8, wherein said neutral moieties are selected from the group consisting of phenethyl, benzyl, phenylpropyl, (R) alpha-methylbenzyl, (S) alpha-methylbenzyl, methoxyphenethyl, and chlorophenethyl.
- 10. The polycationic agent of claim 1, wherein R₁ and R₃ are both hydrogen for at least one monomer.
 - 11. The polycationic agent of claim 10 wherein n is 36.
 - 12. The polycationic agent of claim 10, wherein n is 24.
 - 13. The polycationic agent of claim 10, wherein n is 18.
 - 14. The polycationic agent of claim 10, wherein n is 12.
- 15. The polycationic agent of claim 8, wherein Ta and Tc are terminal groups selected from the group consisting of polypeptide, lipid, lipoprotein, vitamin, hormone, polyakylene glycol, and saccharide.
 - 16. A composition comprising:

- (i) a polynucleotide; and
- (ii) the polycationic agent of claim 1, wherein said polycationic agent is capable of mediating entry of polynucleotides into a cell.
 - 17. A pharmaceutical composition comprising:
 - (i) a pharmaceutically acceptable carrier;
 - (ii) a therapeutically effective amount of polynucleotides; and
- (iii) an amount effective to neutralize said polynucleotides of the polycationic agent of claim 1 wherein said polycationic agent is capable of mediating entry of polynucleotides into a cell.
- 18. A method of complexing polynucleotides with a polycationic agent comprising:
 - (i) providing a polynucleotide; and
- (ii) contacting said polynucleotide with the polycationic agent of claim 1, wherein said polycationic agent is capable of mediating entry of polynucleotides into a cell.
- 19. A method of condensing polynucleotides, said method comprising: contacting a polynucleotide with a condensing amount of the polycationic agent of claim 1, wherein said condensing amount is an amount of polycationic agent sufficient to reduce the size of said polynucleotide.
- 20. A method of inhibiting serum degradation of polynucleotides, said method comprising contacting a polynucleotide with the polycationic agent of claim 1 wherein said polycationic agent is present in an amount effective to inhibit serum degradation by at least 10 minutes.
 - 21. A composition comprising:
 - (i) a lipoprotein;
 - (ii) a polynucleotide binding molecule; and
 - (iii) a polynucleotide,

wherein said composition is capable of increasing the frequency of polynucleotide uptake into a cell.

- 22. The composition of claim 21, wherein the lipoprotein is selected from the group consisting of high density lipoprotein, intermediate density lipoprotein, low density lipoprotein, and very low density lipoprotein.
- 23. The composition of claim 21, wherein the lipoprotein a mutant, fragment or fusion of the protein selected from the group consisting of high density lipoprotein, intermediate density lipoprotein, low density lipoprotein, and very low density lipoprotein.
- 24. The composition of claim 21, wherein the lipoprotein is acetylated low density lipoprotein.
 - 25. A pharmaceutical composition comprising
 - (a) a therapeutically effective amount of a polynucleotide;
- (b) a polynucleotide binding molecule in an amount effective to neutralize the negative charge of said polynucleotide; and
 - (c) a therapeutically effective amount of lipoprotein.
- 26. The pharmaceutical composition of claim 26, wherein said polynucleotide is a polycationic agent.
- 27. A method of producing a composition for facilitating entry of a polynucleotide into a cell said method comprising:
 - (i) providing a polynucleotide
- (ii) providing a polynucleotide binding molecule in an amount effective to neutralize said polynucleotide;
- (ii) contacting said polynucleotide with said polynucleotide binding molecule to form a complex;
 - (iii) providing a lipoprotein; then
 - (iv) contacting the complex with said lipoprotein.